GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
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	primer; hybridisation; illular adhesion disorder;		Murine neokine-1. Murine CXC chemokir Full-length murine Skin cell protein, Macaque neokine-1. Revised partial rodent che Rat neokine-1 (re Revised partial protein, muKS1 fragment (re Skin cell protein, muKS1 fragment (re Mouse muKS1 protein Skin cell protein, Human colon cancer Partial alternativ Murine skin cell protein, Peptide #695 enco Neutrophil chemota Macrophage derived Murine macrophage Gro-beta/MIP-2-alp Human macrophage i Human macrophage i Human mcrophage i Human growth related onc Human chemokine GR Human protein sequ Melanoma growth st Growth related onc Human chemokine GR

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30-JUL-1998;
11-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes secreted and transmembrane polypeptides and their polynucleotides. The nucleotide sequences are useful as sources of probes, primers, for chromosome mapping, and for generation of antisense sequences. They can also be used to create transgenic animals. The proteins can be used to treat a variety of diseases and disorders, depending on their function. Diseases that may be treated include blood coagulation disorders, cancers and cellular adhesion disorders. They may also be used to raise antibodies. AAZ33891 to AAZ34338, and AAY41685 to AAY41774 represent polynucleotide and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New secreted and transmembrane polypeptides and their polynucleotides, useful for treating blood coagulation disorders, cancers and cellular adhesion disorders.
                                                                                                                                                                                                                                                        Chemokine; CXC;
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                                                                                                                                                                                                                                       Tim-1; inflammation; heart attack; stroke; infection;
on; burns; frostbite; corrosive chemical; gene therapy.
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26-JUL-1999
26-JUC-1999
30-NOV-1999
02-DEC-1999
02-DEC-1999
16-DEC-1999
16-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This is the amino acid sequence for the Tim-1 CXC chemokine. The Tim-1 gene can be used to design therapeutic tools for treating inflammation due to stimuli such as heart attacks and stroke, infection, physical trauma, UV or ionizing radiation, burns, frostbite or corrosive chemicals. The Tim-1 gene and subgenomic polynucleotides can be used
                                                                                                                                                                                                                                                                                                                                    WO200053756-A2
                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                expressed sequence tag; detection; cancer.
 Ashkenazi AJ, Baker KP,
Ferrara N, Filvaroff E,
                                                                                                                                                                                                                                                  08-MAR-1999;
12-MAR-1999;
                                                                                                                                                                                                                                                                                                             14-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                          Human; secreted protein; transmembrane protein; PRO;
                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO273 (UNQ240) protein sequence SEQ ID NO:370.
                                                                                                                                                                                                                                                                                                                                                                                                                                          08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB44295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAB44295 standard; Protein; 111 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 43; 46pp; English.
                                     (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                      18-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A new human CXC chemokine, Tim-1, useful for treating inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB; AAX89708
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PHCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           phceekmviittksvsryrggehclhpklgstkrfikwynawnekrrvyee 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MSLLPRRAPPVSMRLLAAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKY 60
                                                        ; 2000WO-US00219.
; 2000WO-US00277.
; 2000WO-US00376.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                     2000WO-US04341
                                                                                                                                                      99US-0145698.
99US-0162506.
99WO-US28313.
                                                                                               99WO-US31274
                                                                                                         99WO-US30095.
99WO-US31243.
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99WO-US28565
                                                                                                                                                                                                                                                             99WO-US05028
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                                                                                                                                                                                                    99US-0134287
                                                                                                                                                                                                                            99US-0130232
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 Botstein D,
Fong S, Gao
Desnoyers L,
W, Gerber H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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 Eaton DL;
Gerritsen ME;
                                                                                                                                                                                                                                                                                                                                                                                           EST;
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                                                                                                                                                                                    haemostatic; antithyroid; antidiabetic; nootropic; neuroprotective; antianaemic; hepatotropic; virucide; antipascoriatic; antiallergic; antiasthmatic; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; spondyloarthropathy; systemic sclerosis; sarcoidosis; idiopathic inflammatory myopathy; Sjogren's syndrome; thyroiditis; systemic vasculitis; autoimmune haemolytic anaemia; diabetes mellitus; autoimmune thrombocytopaenia; immune-mediated renal disease; demyelinating disease; hepatobiliary disease; Whipple's disease; inflammatory bowel disease; gluten-sensitive enteropathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC78458 to AAC78599 represent polynucleotide and EST (expressed sequence tag) sequences which encode secreted or transmembrane PRO polypeptides. The PRO polynucleotides and polypeptides have cytostatic activity The polynucleotides and polypeptides can be used for detecting the presence of PRO polypeptides in samples, for linking bioactive molecules to cells and for modulating biological activities of cells, using the polypeptides for specific targeting. The polypeptide targeting can be used to kill the target cells, e.g. for the treatment of cancers. The polypeptide pairs provide specific targeting of bioactive molecules to cells AAC78600 to AAC78987 represent PCR primers and probes used in the isolation of the PRO polynucleotide sequences.
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Kljavin
                                                                                                                                                        autoimmunde disease; immune-mediated skin disease; immunological disease; transplantation associated
                                                                                                                                         graft rejection;
                                                                                                                                                                                                                                                                                                                                                      Human; immune related disease; diagnosis; antiinflammatory; cardiant; dermatological; antiarthritic; antirheumatic; immunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                          Human PR$0273 protein UNQ240 SEQ ID NO:46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB33423
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel PR\!\!\!\!/ polypeptides and polynucleotides used in detection methods, to target bioactive molecules to specific cells, and to modulate cellular activities -
                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                              29-JAN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 phceekmviittksvsryrgqehclhpklqstkrfikwynawnekrrvyee 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 PHCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4
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DB; AAC78551.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 587; DB 21; $imilarity • 100.0%; Pred. No. 2.3e-60; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;<u>₽</u>₹
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; Protein; 111 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fig 149; 636pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               111 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Godowski PJ,
Kuo SS, Naj
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stewart TA,
                                                                                                                                         graft-versus-host-disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PJ, Grimaldi CJ,
Napier MA, Pan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tumas D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CJ, Gurney AL,
an J, Paoni NF,
Williams PM, Wo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21;
                                                                                                                                                          allergic disease;
disease;
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Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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02-MAR-2000; 2000WO-US05841

14-SEP-2000. WO200053758-A2.

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05-OCT-1999;
29-OCT-1999;
29-NOV-1999;
30-NOV-1999;
30-NOV-1999;
01-DEC-1999;
01-DEC-1999;
02-DEC-1999;
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28-JUL-1999;
01-SEP-1999;
08-SEP-1999;
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20-APR-1999;
28-APR-1999;
04-MAY-1999;
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20-DEC-1999;
30-DEC-1999;
05-JAN-2000;
06-JAN-2000;
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15-SEP-1999;
15-SEP-1999;
anaemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus immune-mediated renal disease, demyelinating diseases of the central and peripheral nervous systems, hepatobiliary diseases, inflammatory bowel disease, jutten-sensitive enteropathy and whipple's disease, autoimmune or immune-mediated skin diseases, allergic disease, immunological diseases of the lung, and transplantation associated diseases including graft rejection and graft-versus-host-disease.

AAC58397 to AAC58578 represent PCR primers and hybridisation probes on the isolation of human PRO sequences. AAC58579 to AAC58642 and
                                                                                                                                                      The present invention describes sixty four human PRO proteins which can be used in the treatment of immune related diseases. The human PRO proteins, anti-PRO antibodies, agonists and antagonists are useful for treating and diagnosing immune related disorders. The disorders are selected from systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis, spondyloarthropathies,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-DEC
                                                                                                                                                                                                                                                                                  Sixty four PRO polypeptides, useful in the diagnosis and treatment of immune related disorders, e.g. systemic lupus erythematosis, rheumatoid arthritis, osteoarthritis, thyroiditis and diabetes mellitus -
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10-MAR-1999;
                                                                                                                                                                                                                                                                                                                                           WPI; 2000-572271/53.
N-PSDB; AAC58588.
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20-JUL-1999
                                                                                                                            systemic sclerosis, idiopathic inflammatory myopathies, Sjogren's syndrome, systemic vasculitis, sarcoidosis, autoimmune haemolytic
                                                                                                                                                                                                                                                         Claim 33; Fig 20; 309pp; English.
                                                                                                                                                                                                                                                                                                                                                                                    Stewart TA,
                                                                                                                                                                                                                                                                                                                                                                                                 Kabakoff RC,
                                                                                                                                                                                                                                                                                                                                                                                                              Ashkenazi AJ,
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11-FEB-2000;
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                                                                                                                                                                                                                                                                                                                                                                                  Tumas D,
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2000WO-US00277.
2000WO-US00376.
2000WO-US03565.
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2000WO-US04414
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                                                                                                                                                                                                                                                                                                                                                                                                  Lu Y,
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99WO-US28565.
99WO-US30095.
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99US-0144758.
99US-0145698.
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99US-0123618.
99US-0123957.
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99WO-US31274
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99WO-US20111
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99US-0132371
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Watanabe CK, Wood WI, Yan M;
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D, Shelton DL,
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DL, Smith
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     AAB88478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                              which encode human secretory or membrane proteins represented by AAB88317 - AAB88419. Included in the invention are primers AAB88317 - AAF64232 - AAF62235 which are used to isolate the CDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA sequences, which can be used in vaccines. The polynucleotide sequences can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and diagnosts of diseases associated with inappropriate secretory protein/membrane protein expression. The nucleic acids and complementary sequences may also be used as DNA probes in diagnostic assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
(e.g. polymerase chain reactions (PCR)) to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB33414 to AAB33477 represent human PRO polynucleotide and protein sequences given in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID 324; 609pp + CD ROM; English.
                                                                                                                                                                                                                                                                                                                                   gene
                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-093989/11.
N-PSDB; AAF93905.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Ota T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; secretory protein; membrane protein; vaccine; gene therapy;
rheumatoid arthritis; diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human membrane or secretory protein clone PSEC0212
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB88478 standard; Protein; 111 AA
                                                                                                                                                                                                                                                                  This invention relates to nucleic acid sequences AAF93744 - AAF93916
                                                                                                                                                                                                                                                                                                                                 Nucleic acids encoding secretory proteins/membrane proteins, useful in gene therapy or as candidate target molecules in drug development -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HELI-) HELIX RES INST.
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2000JP-0183766.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Kawai Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sugiyama
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Hayashi K;
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RESULT
AAW29291
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 111; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human chemokine alpha-2; CKalpha-2; treatment; prevention; tumour; leukaemia; T-cell mediated autoimmune disease; parasitic infection; psoriasis; asthma; allergy; haematopoiesis regulation; growth factor stimulation; angiogenesis inhibition;
The present sequence is human chemokine alpha-2 (CKalpha-2),
                            Claim 14; Pages 66-67;
                                                                                                                        WPI; 1997-480216/44.
                                                                                                                                                   Gentz RL,
                                                                                                                                                                                                         19-MAR-1996;
                                                                                                                                                                                                                                   19-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    wound healing promotion; antagonist; rheumatoid arthritis; autoimmune disease; inflammatory disease; infective disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human chemokine alpha-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  identify modulators (agonists and antagonists) of expression and activity. The antibodies and antagonists may also be used as therapeutic agents to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of the polypeptides in samples (e.g. by enzyme linked immunosorbant assay (ELISA). Examples of diseases which may be treated include rheumatoid
                                                                                                                                                                                                                                                              25-SEP-1997
                                                                                                                                                                                                                                                                                                                              Peptide
                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW29291 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           arthritis and diabetes.
                                                                 Novel human chemokine alpha 2 and DNA encoding it - useful treatment of leukaemia, tumours, autoimmune disease, wound
                                                                                                                                                                              (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                       WO9735010-A1
                                                                                                                                                                                                                                                                                                                                                                        Key
                                                                                                                                                                                                                                                                                                                                                                                                                                        allergic reaction; prostaglandin-independent
                                                                                                                                                                                                                                                                                                                                                         Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MSLLPRRAPPVSMRLLAAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKY 60
                                                                                                                                                                                                                                                                                                                                                                                                                           marrow failure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PHCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     msllprrappvsmrllaaallllllalytarvdgskckcsrkgpkirysdvkklemkpky 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 phceekmviittksvsryrgqehclhpklqstkrfikwynawnekrrvyee 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT86316.
                                                                                                                                                   Li H,
                                                       chronic infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                111 AA;
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                                                                                                                                                                                                        96US-0013653.
                                                                                                                                                                                                                                 97WO-US04329
                                                                                                                                                                                                                                                                                                                 /label= sig_peptide
29..111
/label= mat_peptide
                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                   N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein; 111 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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                            English.
                                                       and fibrotic disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 587; DB 22;
Pred. No. 2.3e-60;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                  healing,
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RESULT
AAY31612
THE CCC X STATES AND A STATES A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   QУ
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chemoattractant cytokine; chemokine; CXC motif; conserved cysteine; signal transduction modulation; angiogenesis inhibition; chemoattraction inhibition; cancer; inflammation; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    which can be used to treat and/or prevent tumours, leukaemia, T-cell mediated autoimmune diseases, parasitic infections, psoriasis, asthma and allergy, regulate haematopoiesis, stimulate growth factor activity, inhibit angiogenesis and promote wound
                                                                                                                           Novel nedkine polypeptides and polynucleotides used for regulating and treating proliferative disorders and diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human neokine-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     healing. CKalpha-2 antagonists can be used to treat rheumatoid arthritis, autoimmune and chronic and acute inflammatory and infective diseases, allergic reactions, prostaglandin-independent
                                                                      Disclosure; Fig 1; 123pp; English.
                                                                                                                                                                                                             WPI; 1999-494271/41.
N-PSDB; AAZ08962.
                                                                                                                                                                                                                                                                                              Barnes TM,
                                                                                                                                                                                                                                                                                                                                                                                                      10-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W09940104-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     post-transplantation organ rejection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY31612
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                                                                                                                                                                                                                                                                                                                                              (MILL-) MILLENNIUM BIOTHERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-AUG-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                110;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    $imilarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bone marrow failure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            111 AA;
                                                                                                                                                                                                                                                                                           MacKay C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                      98US-0023664
                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-US02943
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23..99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "Conserved Cys at 25,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "Human mature neokine-1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /label= CXC_motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein; 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99.1%;
99.1%;
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Pred. No. 8.6e-60;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    51 and 72"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 111;
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This sequence represents human neokine-1. Human sequence tags (ESTs), potentially encoding a che

chemokine,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cysteines separated by one non-conserved amino acid residue; however, the neckines show some atypical features which distinguish them from previously characterised CXC chemokines e.g., the presence of approximately 5 residues between the third and fourth conserved cysteines. The neckines have a variety of functions, including modulation of signal transduction, inhibition of angiogenesis, regulation of inflammation and inhibition of chemoattraction. Neckines may therefore be useful in the treatment of various proliferative disorders or diseases: several types of cancer, inflammation, psoriasis, and immune rejection following skin graft.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       used to assemble two non-overlapping contigs. To establish that these two contigs derived from the same transcript, two primers were designed to amplify the region between these contigs. These primers were hil2/27f (AAX08966) and hil2/526r (AAX08967). To obtain a full open reading frame (ORF), the 5' end of the single contig was generated via the RACE (rapid amplification of cDNA ends) procedure, using primer til2racel (AAZ08968). The neokines are a novel family of chemoattractant cytokines (chemokines) and are ligands for previously identified putative G protein-coupled receptor termed RDC1 (also called the neokine receptor). The neokines are members of the non-ELR-CXC subfamily of chemokines characterised by the absence of an ELR motif and the presence of a CXC signature motif. The CXC motif encompasses 4 highly conserved cysteine residues, with the first two
                                                                                                                                                                                                                                       wound healing promotion; antagonist; rheumatoid arthritis; autoimmune disease; inflammatory disease; infective disease; allergic reaction; prostaglandin-independent fever;
                      WO9735010-A1
                                                                               Peptide
                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                               Human chemokine alpha-2; CKalpha-2; treatment; prevention; tumour; leukaemia; T-cell mediated autoimmune disease; parasitic infection psoriasis; asthma; allergy; haematopoiesis regulation; growth factor stimulation; angiogenesis inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                     Human chemokine alpha-2
                                                                                                                                                                                                                                                                                                                                                                                                                                               14-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW29292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW29292 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and kidney transplantation. Neokines, their associated polynucleotides, homologues and antibodies can be used in screening for drugs or compounds which modulate their activity and therefore may be important for the development of new therapeutics.
                                                                                                                      Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 ksvsryrgqehclhpklqstkrfikwynawnekrrvyee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  73 KSVSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13 MRLLAAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITT 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8
                                                                                                                                                                                                                        marrow failure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mrllaaalllllalytarvdgskckcsrkgpkirysdvkklemkpkyphceekmviitt
                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                 /label= sig_peptide
17..99
                                                                                                                                       Location/Qualifiers
                                                            /label= mat_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     89.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 527; DB 20;
Pred. No. 1.8e-53;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 99;
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                                                                                                                                                                                                                                                                                                                                             infection;
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RESULT
AAY76089
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gentz RL,
                                                                                                                                                                           angiogenesis; tumour vascularisation; growth disorder;
developmental disorder; skin wound; hair follicle disorder;
                                                                                                                                                                                                                  Skin; dermal papilla; keratinocyte; neonatal foreskin fibroblast; embryonic skin cell; keratinocyte stem cell; transit amplifying cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               healing. CKalpha-2 antagonists can be used to treat rheumatoid arthritis, autoimmune and chronic and acute inflammatory and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          which can be used to treat and/or prevent tumours, leukaemia, T-cell mediated autolmmune diseases, parasitic infections, psoriasis, asthma and allergy, regulate haematopoiesis, stimulate growth factor activity, inhibit angiogenesis and promote wound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 14; Pages 66-67; 82pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel human chemokine alpha 2 and DNA encoding it - useful treatment of leukaemia, tumours, autoimmune disease, wound psoriasis, chronic infections and fibrotic disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-MAR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-MAR-1997;
              09-NOV-1998;
                         29-APR-1998;
                                                    29-APR-1999;
                                                                                 04-NOV-1999
                                                                                                           W09955865-A1
                                                                                                                                                                anti-inflammatory;
                                                                                                                                                                                                      secreted; transmembrane; inflammation; cancer; neurological disease;
                                                                                                                                                                                                                                                             Human CXC chemokine homologue huKS1, SEQ ID NO:344
                                                                                                                                                                                                                                                                                        27-MAR-2000
                                                                                                                                                                                                                                                                                                                   AAY76089;
                                                                                                                                                                                                                                                                                                                                             AAY76089 standard; Protein; 95 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is human chemokine alpha-2 (CKalpha-2),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HUMA-) HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-SEP-1997
                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fever and bone marrow failure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Infective diseases, allergic reactions, prostaglandin-independent
                                                                                                                                                                                                                                                                                                                                                                                                                61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13 MRLLAAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITT
                                                                                                                                                                                                                                                                                                                                                                                                                                          73 KSVSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  \vdash
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DB; AAT86316.
                                                                                                                                                                                                                                                                                                                                                                                                              mrllaaallllllalytarvdgskckcsrkgpkirysdvkklemkpkyphceekmviitt 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENOME SCI INC.
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                         98US-0069726
              98US-0188930
                                                      99WO-NZ00051.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ni J,
                                                                                                                                                                 cytostatic; neuroprotective; vulnery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 522; DB 18; Pred. No. 6.7e-53;
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                                                                                                                                                                                                                                                                                                                                                                                                                 99
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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer and neurological diseases. The proteins may be used to stimulate the growth and motility of keratinocytes, to inhibit the growth of cancer cells, to modulate angiogenesis and tumour vascularisation, to modulate skin inflammation, to modulate epithelial cell growth and to inhibit binding of HIV-1 to leukocytes. The invention may also be used to treat growth and developmential defects, skin wounds and hair follicle disorders. Sequences AAY75942-Y76123 represent polypeptides encoded by cDNA sequences AAY75942-Y76123 represent polypeptides encoded types. Sequences AAY75942-Y75947, AAY76020-Y76021, AAY7604-Y76104 and AAY76119 are proteins with an N-terminal signal sequence, indicating that they are secreted. Sequences AAY75986-Y75989, AAY76061-Y76071, AAY76106-Y76109 and AAY76121-Y76122 are proteins with one or more
                                                                                                                                          WO200148192-A1
                                                                                                                                                                                                cell trafficking;
                                                                                                                                                                                                                                                                                                                                               AAE05371 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; Page 202; 235pp; English.
                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                              Human; cytostatic; antiinflammatory; immunoregulatory; tissue integrity;
wound healing; immune response; vaccine; cancer; asthma; allergy;
                                                                                                                                                                                                                                                          Human huKS1 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
            (GENE-) GENESIS RES & DEV CORP LTD
                                        28-NOV-2000;
                                                    23-DEC-1999;
                                                                                  21-DEC-2000; 2000WO-NZ00256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         putative transmembrane domains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to novel nucleic acid sequences derived from rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel polynucleotides useful for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Strachan L,
                                                                                                                                                                                                                                                                                        12-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 including wounds and cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GENE-) GENESIS RES
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                                                                                                                                                                                                                                                                                                                                                                                                                     AAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITTKSVS 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                RYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000-072177/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            papilla, human keratinocytes and neonatal foreskin fibroblasts, use embryonic skin, keratinocyte stem cells and transit amplifying Polypeptides of the invention may be used to treat inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                     2000US-0724864.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sleeman M,
                                                                                                                                                                                                                                                                                      (first entry)
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                                                    99US-0171678
                                                                                                                                                                                                  therapy; huKS1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ę,
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                                                                                                                                                                                                                                                                                                                                               95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 509; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                               A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               treatment of various conditions
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2e-51;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cc antibodies, to isolate corresponding interacting proteins or other compounds, to quantitatively determine levels of interacting proteins or other compounds, and as therapeutic target in a whole range of disease cc states. Compositions comprising the novel proteins of the invention are useful for treating mammalian disorders. Polynucleotides of the invention are cuseful in genome and physical mapping, in positional cloning of cc genes, to tag or identify an organism or its reproductive material (as non-disruptive tags for marking organisms), and for the diagnosis and cc treatment of mammalian diseases which is the consequence of inappropriate expression of kinase genes. They are useful for promoting immune response cc as part of a vaccine or anti-cancer treatment, as immunoregulatory and anti-inflammatory molecule, as citagnostic for specific types of cancer and for development of an confidence or enhance the activity of the soluble molecule that binds proteins of the invention, for tissue and neural regeneration, to promote or block call trafficking, and as anti-inflammatory and/or vaccine adjuvant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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23-NOV-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The patcht discloses novel polynucleotides and their corresponding proteins which play a major role in induction of growth, cell migration and proliferation, cell-cell interaction and the differentiation of tissue-specific cells. These proteins are important in the maintenance of tissue integrity and thus are important in wound healing. They are useful in various assays to determine the biological activity, to raise
                                                                                                                                                                                            Skin cel
                                                                                                                                                                                                                             08-MAR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel is blated polypeptide useful to isolate corresponding interacting proteins or other compounds, to quantitatively determine levels of
                                 WO200069884-A2
                                                                  Homo sap
                                                                                                                    nootropic; neuroprotective; vulnerary; in
keratinocyte growth stimulation; cancer;
                                                                                                                                                         Human; skin cell; cytostatic; antiinflammatory; anti-HIV;
                                                                                                                                                                                                                                                                                                     AAB56028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example
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                                                                                                         inflamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence is human huKS1 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            interacting proteins or other compounds, and as therapeutic target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17
                                                                                                                                                                                                                                                                                                                                        11
                                                                                                                                                                                                                                                                                                                                                                                      61 ryrqqehclhpklqstkrfikwynawnekrrvyee
                                                                                                                                                                                                                                                                                                                                                                                                                           77 RYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                          95;
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                                                                    lens.
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                                                                                                                                                                                                                                                                                                   standard; Protein;
                                                                                                                                                                                            protein,
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                                                                                                     neurological disease
                                                                                                                                                                                          SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                       immunomodulatory; vaccine;
r; angiogenesis inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2e-51;
                                                                                                                                                                                                                                                                                                                                                                                            95
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RESULT
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is a polypeptide which is expressed in mammalian skin cells. The polypeptide is useful for stimulating keratinocyte growth and motility, inhibiting the growth of cancer cells, modulating angiogenesis, inhibiting angiogenesis and vascularisation of tumours, modulating skin inflammation, stimulating the growth of epithelial cells, inhibiting the binding of human immunodeficiency virus (HIV)-1 to leukocytes, and treating inflammatory disease, cancer and neurological diseases. The polynucleotide can be used as a marker, in the identification of genetic disorders, and for the design of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated polynucleotide used in the identification of genetic disorders and encoding polypeptides used for treating inflammatory disease, cancer and neurological diseases - \,
                                                                                                                                                                                                                                                                                                                                                                                                   signal transduction modulation; angiogenesis inhibition; chemoattraction inhibition; cancer; inflammation; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chemoattractant cytokine; chemokine; CXC motif; conserved cysteine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY31613 standard; Protein; 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; Page 270; 352pp; English.
                              WO9940104-A1
                                                                                                                                                                                                                                             Peptide
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                                                                                                                                                                                                                                                                                                                                                                                post-transplantation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17 AAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITTKSVS 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                            /note=
                                                                                                                                                                                                                 /label= Signal_peptide
                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                          /label= CXC_motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for examining expression patterns.
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100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                organ
                                                                                                                                                               "Murine mature neokine-1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Onrust R,
                                                                                                                                                                                                                                                                                                                                                                                rejection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 509;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                              Cys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sleeman M,
                                                                                at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2e-51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 22;
                                                                                25,
                                                                                   27,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 95;
                                                                                   51
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                                                                                and
                                                                                   72"
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MRLLAAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITT

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ksmsryrgqehclhpklqstkrfikwynawnekrrvyee KSVSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111

99

AAY76085

13

AAY76085 standard; Protein;

99

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secreted; transmembrane; inflammation; cancer; neurological disease; angiogenesis; tumour vascularisation; growth disorder; developmental disorder; skin wound; hair follicle disorder;

anti-inflammatory; cytostatic; neuroprotective; vulnery

Skin; dermal papilla; keratinocyte; neonatal foreskin fibroblast; embryonic skin cell; keratinocyte stem cell; transit amplifying c

cell;

Murine CXC chemokine homologue muKS1, SEQ ID NO:340.

27-MAR-2000 AAY76085;

(first entry)

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Matches
                                  Query Match
Best Local
                                                                                                                                                                                                                                                                               cysteines. The mockines have a variety of functions, including modulation of signal transduction, inhibition of angiogenesis, regulation of inflammation and inhibition of chemoattraction. Neokines may therefore be useful in the treatment of various proliferative disorders or diseases: several types of cancer, inflammation, psoriasis, and immune rejection following skin graft and kidney transplantation. Neokines, their associated polynucleotides, homologues and antibodies can be used in screening for drugs or compounds which modulate their activity and therefore may be important for the description.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 previously identified putative G protein-coupled receptor termed RDC1 (also called the neckine receptor). The neckines are members of the non-ELR-CXC subfamily of chemokines characterised by the absence of an ELR motif and the presence of a CXC signature motif. The CXC motif encompasses 4 highly conserved overtice and the presence of a call of the call
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cysteines separated by one non-conserved amino acid residue; how the neokines show some atypical features which distinguish them previously characterised CXC chemokines e.g., the presence of approximately 5 residues between the third and fourth conserved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ELR motif and the presence of a CXC signature motif. The CXC motif encompasses 4 highly conserved cysteine residues, with the first two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel neokine polypeptides and polynucleotides used and treating proliferative disorders and diseases
                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence represents murine neokine-1. The neokines are a novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Fig 2; 123pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Barnes TM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-FEB-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-AUG-1999
                                                                                                                                                                                                                                                        the development of new therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1999-494271/41.
                                      Similarity
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   Conservative
                                                                                                                                                                                      ΑĀ,
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                                  86.2%;
94.9%;
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••
                                  Score 506; DB 20; Pred. No. 4.7e-51;
   Mismatches
2;
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Best Local :
                    Skin; dermal papilla; keratinocyte; neonatal foreskin fibroblast; embryonic skin cell; keratinocyte stem cell; transit amplifying cell secreted; transmembrane; inflammation; cancer; neurological disease; angiogenesis; tumour vascularisation; growth disorder; developmental disorder; skin wound; hair follicle disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer and neurological diseases. The proteins may be used to stimulate the growth and motility of keratinocytes, to inhibit the growth of cancer cells, to modulate angiogenesis and tumour vascularisation, to modulate skin inflammation, to modulate epithelial cell growth and to inhibit binding of HIV-1 to leukocytes. The invention may also be used to treat growth and developmental defects, skin wounds and hair follicle disorders. Sequences AAV75942-Y76123 represent polypeptides encoded by cDNA sequences AAV75942-Y7613 represent polypeptides encoded by cDNA sequences derived from several mouse, rat or human skin cell types. Sequences AAV75942-Y7547, AAY76021, AAY76094-Y76104 and AAX76119 are proteins with an N-terminal signal sequence, indicating that they are secreted. Sequences AAY75986-Y75989, AAY76061-Y76071, AAY76106-Y76109 and AAX76111-Y76122 are proteins with one or more notification to the sequence of the 
                                                                                                                                                                                    Full-length murine CXC chemokine homologue muKS1, SEQ ID NO:394.
                                                                                                                                                                                                                                                      27-MAR-2000
                                                                                                                                                                                                                                                                                                                                                           AAY76115 standard; Protein; 99 AA
anti-inflammatory; cytostatic; neuroprotective; vulnery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and mouse embryonic skin, keratinocyte stem cells and transit amplifying cells. Polypeptides of the invention may be used to treat inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to novel nucleic acid sequences derived from rat dermal papilla, human keratinocytes and neonatal foreskin fibroblasts.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; Page 200; 235pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB; AAZ61790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-072177/06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    including wounds and cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9955865-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            73 KSVSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polynucleotides useful for the treatment of various conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                86.2%;
94.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Watson JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 506; DB 21;
Pred. No. 4.7e-51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Onrust R,
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     Mouse; skin cell; cytostatic; antiinflammatory; anti-HIV; nootropic; neuroprotective; vulnerary; immunomodulatory; vaccine; keratinocyte growth stimulation; cancer; angiogenesis inhibition; inflammation; neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer and neurological diseases. The proteins may be used to stimulate the growth and mothlity of keratinocytes, to inhibit the growth of cancer cells, to modulate angiogenesis and tumour vascularisation, to modulate skin inflammation, to modulate epithelial cell growth and to inhibit binding of HIV-1 to leukocytes. The invention may also be used to treat growth and developmental defects, skin wounds and hair follicle disorders. Sequences AAV75942-Y76123 represent polypeptides encoded by cDNA sequences derived from several mouse, rat or human skin cell types. Sequences AAV75942-Y75947, AAY76020-Y76021, AAY76094-Y76104 and
                                                                                                 Skin cell protein, SEQ ID NO:
                                                                                                                                     08-MAR-2001
                                                                                                                                                                                                            AAB56024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY76119 are proteins with an N-terminal signal sequence, indicatin that they are secreted. Sequences AAY75986-Y75989, AAY76061-Y76071, AAY76106-Y76109 and AAY76121-Y76122 are proteins with one or more
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to novel nucleic acid sequences derived from rat dermal papilla, human keratinocytes and neonatal foreskin fibroblasts, and mouse embryonic skin, keratinocyte stem cells and transit amplifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cells. Polypeptides of the invention may be used to treat inflammation cancer and neurological diseases. The proteins may be used to stimulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-APR-1998;
09-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 223; 235pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel polynucleotides useful for the treatment of various conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Strachan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
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                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               transmembrane domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           wounds and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                     (first entry)
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                                                                                                                                                                                                        Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 506; DB 21;
Pred. No. 4.7e-51;
                                                                                                 340.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                          The present sequence is a polypeptide which is expressed in mammalian skin cells. The polypeptide is useful for stimulating keratinocyte growth and motility, inhibiting the growth of cancer cells, modulating angiogenesis, inhibiting angiogenesis and vascularisation of tumours, modulating skin inflammation, stimulating the growth of epithelial cells, inhibiting the binding of human immunodeficiency virus (HIV)-1 to leukocytes, and treating inflammatory disease, cancer and neurological diseases. The polynucleotide can be used as a marker, in the identification of genetic disorders, and for the design of oligonucleotides for examining expression patterns.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated polynucleotide used in the identification of genetic disorders and encoding polypeptides used for treating inflammatory disease, cancer and neurological diseases \dot{}
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; Page 268; 352pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-007495/01.
N-PSDB; AAC99723.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Watson JD, Strachan L,
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                                                                                                                                                                                                                                                                 Sequence
                                                                     94;
                                                                                                                                                                                                                                                                    99 AA;
                                                                                                                                                                                             Conservative
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                                                                                                                                                                                           86.2%; score 506; DB 22; Length 99; 94.9%; pred. No. 4.7e-51; tive 3; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kumble KD,
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Result
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Maximum Match 100%
Listing first 45 summaries
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         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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DB seq length: 2000000000
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Copyright (c) 1993 - 2000 Comp
            /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
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US-08-825-556A-3
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US-09-188-930-345
US-09-188-930-157
US-09-188-930-157
US-08-825-556A-4
US-08-825-556A-4
US-08-825-556A-4
US-08-352-324A-4
US-08-352-324A-4
US-08-352-324A-4
US-08-352-324A-7
US-08-825-324A-7
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PCT-US95-16144-7
US-08-825-324A-7
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US-08-825-324A-7
US-08-825-324A-7
US-08-825-324A-5
US-08-852-324A-5
US-08-852-324A-5
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US-08-862-607-5
US-08-862-607-5
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US-08-862-607-5
US-08-468-819-7
US-08-862-607-5
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US-08-862-607-5
US-08-468-819-7
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Query Match
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Matches 110; Conserv

Conservative

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99.1%; 99.1%;

Score 582; DB 2; Pred. No. 8.6e-64; Mismatches

Length 111; Indels

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RESULT 1 US-08-825-556A-2 Sequence 2, Apperent No. 991 Fatent No. 991 GENERAL INFORMATION FOR APPLICANT APPLICANT APPLICANT APPLICANT TITLE OF IN NUMBER OF SET COUNTRY COMPUTER RESELY COUNTRY COMPUTER REFERENCE CURRENT APPLICANT FILING DA APPLICATI FILING DA ANAME: St RESERENCE TELECOMMUNI TELEPHONE TELECOMMUNI TELEPHO	28 99.5 30 99.5 31 32 94.5 33 34 88.5 35 88.5 36 88.5 37 88.5 38 88.5 41 41 87.5 87.5 87.5
1 25-556A-2 25-55A-2	17.0 73 2 US-08-436-420-42 Sequence 42 5 17.0 74 3 US-08-679-493A-147 Sequence 3, 7 16.5 96 4 US-08-679-493A-147 Sequence 14 16.3 78 2 US-08-436-420-38 Sequence 14 16.1 126 1 US-07-624-742-1 Sequence 3, 15.9 72 2 US-08-436-420-43 Sequence 3, 15.1 72 3 US-08-846-966-1 Sequence 4, 15.1 72 3 US-08-557-142-1 Sequence 1, 15.1 72 5 PCT-US94-06364-1 Sequence 1, 15.1 72 1 US-08-436-420-41 Sequence 1, 15.1 72 2 US-08-436-420-41 Sequence 1, 15.1 72 1 US-07-7792-988-3 14.9 73 1 US-07-778-413E-19 Sequence 3, 14.9 73 1 US-08-346-102-19 Sequence 1, 14.9 73 1 US-08-346-20-27 14.9 73 3 US-08-846-966-3 Sequence 2, 14.9 73 3 US-08-846-966-3 Sequence 3, 14.9 73 3 US-08-846-966-3 Sequence 3, 14.9 73 3 US-08-846-966-3 Sequence 3, 14.9 73 5 PCT-US94-06284-3 Sequence 3,
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US-09-188-930-344; Sequence 344, Application US/09188930A; Patent No. 6150502
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GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/013,653
FILING DATE: 19-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,668
REFERENCE/DOCKET NUMBER: 1488.08500
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ni, Jian APPLICANT: Gentz, Reiner L.
                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                           61
                                                                                                                                            73 KSVSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111
                                                                                                                                                                                                                   13 MRILAAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITT 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 202-371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/825,556A FILING DATE: 19-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 20005-2934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Washington STATE: DC
                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C STREET: 1100 New York Ave., Suite 600
                                                                                                                                                                                                1 MRLLAAALLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITT 60
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                                                                                                                         KSVSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRFYEE 99
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Li, Haodong
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RESULT 5
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                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 344
LENGTH: 95
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 340, Application US/09188930A Patent No. 6150502
                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Watson, James D.
APPLICANT: Strachan, Lorna
APPLICANT: Strachan, Matthew
APPLICANT: Onrust, Rene
APPLICANT: Onrust, Rene
APPLICANT: Murison, James Greg
APPLICANT: ON INVENTION: Compositions Isolated From Skin Cells
TITLE OF INVENTION: and Methods For Their Use
TILE REFERENCE: 11000.1011c1
CURRENT APPLICATION NUMBER: US/09/188,930A
CURRENT FILING DATE: 1998.11-09
NUMBER OF SEQ ID NOS: 348
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/188,930A CURRENT FILING DATE: 1998-11-09 NUMBER OF SEQ ID NOS: 348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Watson, James D. APPLICANT: Strachan, Lorna APPLICANT: Sleeman, Matthew APPLICANT: Onrust, Rene APPLICANT: Murison, James Greg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Compositions Isolated From Skin Cells TITLE OF INVENTION: and Methods For Their Use FILE REFERENCE: 11000.1011c1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Mouse
                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH: 99
                                                                                61
                                                                                               73 KSVSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111
                                                                                                                                                                                13 MRLLAAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITT 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   77 RYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17 AAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITTKSVS 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 RYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 AAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITTKSVS 60
                                                                                                                                                             MRLLAAALLLLLLALCASRVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIVTT 60
                                                                              KSMSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE
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                                                                                                                                                                                                                                          94;
                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                             86.2%;
94.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                          ω
                                                                                                                                                                                                                                        Score 506; DB 4;
Pred. No. 1.4e-54;
3; Mismatches 2
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                                                                                                                                                                                                                                                                             DB 4; Length 99;
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Sequence 346, Application US/09188930A Patent No. 6150502

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APPLICANT: Strachan, Lorna
APPLICANT: Sleeman, Matthew
APPLICANT: Onrust, Rene
APPLICANT: Onrust, Rene
APPLICANT: Murison, James Greg
TITLE OF INVENTION: Compositions Isolated From Skin Cells
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1011c1
CURRENT APPLICATION NUMBER: US/09/188,930A
CURRENT ETLING DATE: 1998-11-09
NUMBER OF SEQ ID NOS: 348
SOSTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 346
LENGTH: 77
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RESULT 7
US-09-188-930-157
; Sequence 157, Application US/09188930A
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CURRENT FILING DATE: 1998-11-09
NUMBER OF SEQ ID NOS: 348
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 345
LENCTH: 77
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Watson, James D.
APPLICANT: Strachan, Lorna
APPLICANT: Strachan, Matthew
APPLICANT: Onrust, Rene
APPLICANT: Murison, James Greg
TITLE OF INVENTION: Compositions Isolated From Skin Cells
TITLE OF INVENTION: and Methods For Their Use
TITLE REFERENCE: 11000.1011c1
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                                                                                                                                                                  95 FIKWYNAWNEKRRVYEE 111
                                                                                                                                                                                                                                                                                                    Local Similarity
mes 75; Conserv
                                                                                                                        61 FIKWYNAWNEKRRVYEE 77
                                                                                                                                                                                                             1 SKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIVTTKSMSRYRGQEHCLHPKLQSTKR 60
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                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                  72.2%;
97.4%;
                                                                                                                                                                                                                                                                                               Score 424; DB 4; Length 77; Pred. No. 1.1e-44; Indels 2; Mismatches 0; Indels
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1157
LENGTH: 133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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TITLE OF INVENTION: Compositions Isolated From Skin Cells
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1011c1
CURRENT APPLICATION NUMBER: US/09/188,930A
CURRENT FILING DATE: 1998-11-09
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TELEFÁX: 202-371-2540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
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APPLICANT:
APPLICANT:
APPLICANT:
                                                         REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 144
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/013,653
FILING DATE: 19-MAR-1996
ATTORNEY/ACENT INFORMATION:
NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,688
                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION UNMBER: U$908/825,556A
FILING DATE: 19-MAR-1997
                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUN TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Chemokine Alpha 2
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 202-371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 1100 New CITY: Washington
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                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 20005-2934
                                                                                                                                                                                                                                                                                                                                                                                                                      STATE:
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: Strachan, Lorna
: Sleeman, Matthew
: Onrust, Rene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    1100 New York Ave., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Su, Jeffrey Y.
Li, Haodong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gentz, Reiner L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ni, Jian
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63.7%;
                          4:
                                                                                                  1488.0850001
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Pred. No. 5.2e-29;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 4; Length 133;
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                                                                                                                                                                                                                                    Sequence 2, Application US/08476376 Patent No. 6103234
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Taylor, Ethan W. TITLE OF INVENTION: SELENOPROTEINS, CODING SEQUENCES AND METHODS FILE REFERENCE: 55-95
                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 216
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
                                                                        APPLICANT: WOLPE, STEPHEN D.
APPLICANT: CERAMI, ANTHONY
APPLICANT: SHERKY, BARBARA
APPLICANT: TEKAMP-OLSON, PATRICIA A.
TITLE OF INVENTION: MACROPHAGE-DERIVED INFLAMMATORY MEDIATOR
TITLE OF INVENTION: (MIP-2)
NUMBER OF SEQUENCES: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 98 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: peptide
                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 24.48;
Local Similarity 35.18;
                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                        61 PHCTQTEVIATLKD-----GQEVCLNPQAPRLQKIIQ 92
                                                                                                                                                                                                                                                                                                                                                                                             61 PHCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
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      2: DAVID A. JACKSON, ESQ.
411 HACKENSACK AVE, CONTINENTAL PLAZA, 4TH
                                                                                                                                                                                                                                                                                                                                                                                                                                                         FLOOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24.4%; Score 143.5; DB 2 nilarity 35.1%; Pred. No. 2.7e-10; Conservative 19; Mismatches 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 143.5; DB 4
Pred. No. 2.7e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels 13;
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                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                       ANTI-SENSE:
                                                                                                                                                                                                                                                                        MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: 201-487-5800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 08-SEP-1981
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 0 FILING DATE: 22-FEB-1982 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 02-SEP-PRIOR APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 01-SEP-1989 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/285,498
FILING DATE: 03-AUG-1994
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61 HCAQTEVIATLKG-----GQKVCLDPEAPLVQKIIQ 91
                                62 HCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: JACKSON ESQ., DAVID A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 13046 CIP A2D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 0 FILING DATE: 08-SEP-1981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 07 FILING DATE: 02-OCT-1987
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OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 0 FILING DATE: 10-AUG-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/476,376 FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
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                                                                                                        8 APPVSMRLLAAALLLLLLALYTARVDGS-----KCKCSRKGPKIRYSDVKKLEMKPKYP 61
                                                                    APP-TCRLLSAALVLLLLLATNHQATGAVVASELRCQCLKTLPRVDFKNIQSLSVTPPGP 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HACKENSACK
                                                                                                                                                                                                                                                                                                                                              amino acid
                                                                                                                                                                                                                                                                                                                                                                 100 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                        201-343-1684
                                                                                                                                             Conservative
                                                                                                                                                                                                                                                         NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-SEP-1988
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13-JUL-1995
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                                                                                                                                                            23.5%; Score 138; DB 3; Length 100; 33.3%; Pred. No. 1.3e-09;
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                                                                                                                                                                              Sequence 4, Application US/08862607 Patent No. 5844084
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                               GENERAL INFORMATION:
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
APPLICANT: Wilde, Craig G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 4:
APPLICANT: Wilde, Craig G. APPLICANT: Seilhamer, Jeffrey J. APPLICANT: Seilhamer, Jeffrey J. TITLE OF INVENTION: A NOVEL CHEMOKINE EXPRESSED IN TITLE OF INVENTION: INFLAMED ADENOID, ITS PRODUCTION AND USES NUMBER OF SEQUENCES: 9

CORRESPONDENCE ADDRESS:
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ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/352,324A
FILING DATE: 07-DEC-1994
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TITLE OF INVENTION: A NOVEL CHEMOKINE EXPRESSED IN
TITLE OF INVENTION: INFLAMED ADENOID, ITS PRODUCTION AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS: LENGTH: 107 amino acid
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                                                                                                                                                                                                                                                                                                 68 HCAQTEVIATLKN-----GQKACLNPASPMVKKIIE 98
                                                                                                                                                                                                                                                                                                                                      62 HCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                 8 APPVSMRLLAAALLLLLLALYTARVDGS------KCKCSRKGPKIRYSDVKKLEMKPKYP 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    107 amino acids
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Hawkins, Phillip R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PF-0025 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 128.5; DB Pred. No. 2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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US-08-468-819-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; MOLECULE TYPE: peptide US-08-862-607-4
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                                                                                                                                                                                                                                                                                                                                                        Sequence 6,
Patent No.
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Matches
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CLASSIFICATION: 435
CLASSIFICATION DATA: 435
PRIOR APPLICATION NUMBER: 08/352,32
APPLICATION NUMBER: 08/352,32
APPLICATION UNBER: 08/352,32
APPLICATION NUMBER: 08/352,32
APPLICATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PP-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                             ZIP: 77210
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                     APPLICANT: Strieter, Robert M.
APPLICANT: Polverini, Peter J.
APPLICANT: Kunkel, Steven L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/862,607
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                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee
                                                                                                                                                                                                                                                 APPLICANT: Kunkel, Steven L.
TITLE OF INVENTION: CXC Chemokines as Regulators of
TITLE OF INVENTION: Angiogenesis
                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                    STREET:
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SOFTWARE: FastSEQ Version 1.5
                                                                                                                                  COUNTRY:
                                                                                                                                                      STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            68 HCAQTEVIATLKN-----GQKACLNPASPMVKKIIE 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            62 HCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
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STREET: 31
 APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
32; Conserv
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                                                                                                                                                                  Houston
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                                                                                                                                                                                     P.O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Incyte Pharmaceuticals, Inc. 3174 Porter Drive
                                                                                                                                    SD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IBM Compatible
                                                                                 Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21.9%; Score 128.5; DB 33.3%; Pred. No. 2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08/352,324
US/08/468,819
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 107;
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ATTORNEY/AGENT INFORMATION:

FILING DATE: CONTROL CLASSIFICATION:

Concurrently herewith N: 424

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US-09-203-235-4
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
             TELECOMMUNICATION INFORMATION: TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Wilde, Craig G.
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: A NOVEL CHEMOKINE EXPRESSED IN
TITLE OF INVENTION: INFLAMED ADENOID, ITS PRODUCTION AND USES
                                                                                         APPLICATION NUMBER: 08/352,324
FILING DATE: 07-DEC-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                 ZIP: 94304
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
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                                                            NAME: Luther, Barbara REGISTRATION NUMBER:
                                                                                                                                           APPLICATION NUMBER: 08/862,607 FILING DATE: 23-MAY-1997
                                            REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                  FILING DATE:
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                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/203,235
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Guegler, Karl J.
Hawkins, Phillip R.
415-852-0195
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Best Local Similarity
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                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                           TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/352,324 FILING DATE: 07-DEC-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: INCYTE PHARMACEUTICALS, INC.
TITLE OF INVENTION: A NOVEL CHEMOKINE EXPRESSED IN INFLAMED
TITLE OF INVENTION: ADENOID, ITS PRODUCTION AND USES
NUMBER OF SEQUENCES: 9
                                                                                              MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
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                 Local Similarity
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CLASSIFICATION:
                                                                                                                                              TYPE:
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                                                                                                                TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8 APPVSMRLLAAALLLLLLALYTARVDGS-----KCKCSRKGPKIRYSDVKKLEMKPKYP 61
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    Conservative
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SYSTEM: PC-DOS/MS-DOS
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07-DEC-1995
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21.9%; Score 128.5; DB 5;
33.3%; Pred. No. 2e-08;
"10. Mismatches 35;
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33.3%;
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    Indels 11;
                                Length 107;
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8 APPVSMRLLAAALLLLLLALYTARVDGS-----KCKCSRKGPKIRYSDVKKLEMKPKYP 61 | | : | | | : :: :::| |

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Search completed: January 29, 2002, 16:04:25 Job time: 51 sec

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OM protein - protein search, using sw model
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Maximum Match 100%
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                No.
           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and' is derived by analysis of the total score distribution.
        Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match
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Copyright (c) 1993 - 2000 Compugen Ltd
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RESULT 2 \$346198 cytokine-induced neutro Cytokines: Rattus norve C;Species: Rattus norve C;Date: 20-Feb-1995 #se C;Date: 20-Feb-1995 #se C;Date: 20-Feb-1995 #se C;Accession: \$46198; C: R;Nakagawa, H; Komorii Biochem. J. 301, 545-5; A;Tatle: Identification ences and characterizal A;Reference number: \$4 A;Accession: \$46198 A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-100 <nak 45,="" a40<="" a;cross-references:="" a;reference="" a;title:="" biochem.="" emn="" h;="" ikesue,="" number:="" of="" pharmacol.="" production="" r;nakagawa,="" td=""><td>Query M Best Lo Matches Qy 13 Db 1 Qy 73 Db 61</td><td>RESULT 1 JG0182 chemokine BRAK - hume C;Species: Homo sapis C;Species: Homo sapis C;Date: 23-Jul-1999 4 C;Accession: JG0182 R;Hromas, R.; Broxmey Blochem. Biophys. Res A;Title: Cloning of E A;Reference number: 3 A;Accession: JG0182 A;Status: preliminar; A;Molecule type: mRN/ A;Residues: 1-99 <hrs a;reross-references:="" beta-t<="" c="" c;superfamily:="" td=""><td></td><td>30 31 33 33 33 34 44 44 44 44 44 44 44 44 44</td></hrs></td></nak>	Query M Best Lo Matches Qy 13 Db 1 Qy 73 Db 61	RESULT 1 JG0182 chemokine BRAK - hume C;Species: Homo sapis C;Species: Homo sapis C;Date: 23-Jul-1999 4 C;Accession: JG0182 R;Hromas, R.; Broxmey Blochem. Biophys. Res A;Title: Cloning of E A;Reference number: 3 A;Accession: JG0182 A;Status: preliminar; A;Molecule type: mRN/ A;Residues: 1-99 <hrs a;reross-references:="" beta-t<="" c="" c;superfamily:="" td=""><td></td><td>30 31 33 33 33 34 44 44 44 44 44 44 44 44 44</td></hrs>		30 31 33 33 33 34 44 44 44 44 44 44 44 44 44
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IT 2 ine-induced neutrophil checies: Raftus norvegicus (1) e: 20-Feb-1995 #sequence_ ession: 846198; C48988 agawa, H.; Komorita, N.; em. J. 301, 545-550, 1994 ie: Identification of cyt. and characterization. erence number: 846198; MU ession: 846198; MU tus: preliminary ecule type: mRNA idues: 1-100 <ans 1-100="" 1425-1.="" 45,="" <ans="" a.;="" a48988;="" agawa,="" an="" d2109;="" ecule="" em.="" embl:="" erence="" h.;="" ha;="" idues:="" ikesue,="" inte-="" le:="" mrna="" mu<="" number:="" of="" pharmacol.="" preliminary="" production="" references:="" ss="" td="" tus:="" type:=""><td>Similarity 3; Conser AAALLLLL AAALLLLLL SAYRGQEHCL </td><td>k - human mo sapiens (h-1999 #sequ JG0182 Broxmeyer, hys. Res. Co ing of BRAK, umber: JG018 Ilminary pe: mRNA -99 <hro> -99 <hro> -99 <hro -99="" <hr<="" <hro="" td=""><td></td><td>112.9 112.56 111.66</td></hro></hro></hro></td></ans>	Similarity 3; Conser AAALLLLL AAALLLLLL SAYRGQEHCL	k - human mo sapiens (h-1999 #sequ JG0182 Broxmeyer, hys. Res. Co ing of BRAK, umber: JG018 Ilminary pe: mRNA -99 <hro> -99 <hro> -99 <hro -99="" <hr<="" <hro="" td=""><td></td><td>112.9 112.56 111.66</td></hro></hro></hro>		112.9 112.56 111.66
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induced neutrophil chemoattractant Raitus norvegicus (Norway rat) O-Feb-1995 #sequence_revision 20-F Oon: \$46198; C48988 I, KOMORITA, N.; Shibata, F.; J. 301, 545-550, 1994 Identification of cytokine-induced characterization ice number: \$46198; MUID:94318061 con: \$46198 preliminary e type: mRNA sis: 1-100 <nak> sis: 1-100 <nak> sis: 1-100 <nak> sis: 1-100 <nak 1425-1430,="" 1993="" 45,="" a48988;="" an="" ce="" hatakeyama,="" interlewkin-8-lik="" muid:93228656<="" number:="" of="" pharmacol,="" production="" s.;="" td=""><td>; Score ; Pred. 0; Mis GSKCKCSRK </td><td>ion 2 C.; 703- iverg 16041</td><td>ALIGN</td><td>T27127 T32670 A54188 A26774 146871 B54188 A37034 728087 576941 15927 151685 T25246 S76323 C84251 S69133 JE0337</td></nak></nak></nak></nak>	; Score ; Pred. 0; Mis GSKCKCSRK 	ion 2 C.; 703- iverg 16041	ALIGN	T27127 T32670 A54188 A26774 146871 B54188 A37034 728087 576941 15927 151685 T25246 S76323 C84251 S69133 JE0337
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R;Anisowicz, A.; Bardwell, L.; Sager, R.
Proc. Natl. Acad. Sci. U.S.A. 84, 7188-719
A;Title: Constitutive overexpression of a
A;Reference number: A94184; MUID:88041072
A;Accession: B28414
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R;Tekamp-Olson, P.; Gallegos, C.; Bauer, D.; McClain, J.; Sherry, B.; Fabre, M.; van Dev J. Exp. Med. 172, 911-919, 1990
A;Title: Cloning and characterization of cDNAs for murine macrophage inflammatory protei A;Reference number: JH0200; MUID:90354792
A;Accession: JH0200
                                                                                                                                    growth-regulated protein precursor - Chinese hamster C;Species: Cricetulus griseus (Chinese hamster) C;Date: 30-Jun-1989 #sequence_revision 30-Jun-1989 #C;Accession: B28414
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C;Species: Mus musculus (house mouse)
C;Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 20-Aug-1999
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A;Molecule type: protein
A;Residues: 33-56 <NA2>
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C;Keywords: heparin binding
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A; Molecule type: mRNA
                                                                                                                                                                                                                                          B28414
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         F;1-27/Domain: signal sequence #status predicted <SIG>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: mRNA
A; Residues: 1-100 <TEK>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPVSMRLLAAALLLLLLALYTARVDGS-----KCKCSRKGPKIRYSDVKKLEMKPKYP
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Pred. No. 2.2e-07;
1; Mismatches 31;
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Pred. No. 5.9e-08
                                                                                         7188-7192,
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30-Jun-1989 #text_change
                                                                growth-regulated
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A;Molecule type: mRNA
A;Residues: 56-107 <SPO>
C;Superfamily: beta thromboglobulin
C;Keywords: inflammation
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A; Residues: 1-107 <IID>
A; Residues: 1-107 <IID: g183627 A; Cross-references: GB: M57731; GB: M36964; NID: g183626; PIDN: AAA63182.1; PID: g183627 A; Haskill, S.; Peace, A.; Morris, J.; Sporn, S.A.; Anisowicz, A.; Lee, S.W.; Smith, Proc. Natl. Acad. Sci. U.S.A. 87, 7732-7736, 1990 A; Title: Identification of three related human GRO genes encoding cytokine function: A; Reference number: A38290; MUID: 91017578
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: GB:X53799; NID:g34658; PIDN:CAA37808.1; PID:g34659 R;Iida, N.; Grotendorst, G.R. Mol. Cell. Biol. 10, 5596-5599, 1990 A;Title: Cloning and sequencing of a new gro transcript from activated A;Reference number: A35931; MUID:90377259 A;Accession: A35931
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A;Cross-references: GB:J03560; NID:g191088; PIDN:AAA36985.1; PID:g304509
A;Note: the authors translated the codon CAG for residue 52 as Glu
C;Superfamily: beta-thromboglobulin
C;Superfamily: beta-thromboglobulin
E;1-23/Domain: signal sequence #status predicted <SIG>
                                                                                                                                                   F;1-34/Domain: signal sequence #status predicted <SIG>
F;35-107/Product: macrophage inflammatory protein 2 alpha #status predicted <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: GB:M36820; NID:g183628; PIDN:AAA63183.1; PID:g183629 R;Sporn, S.A.; Elerman, D.F.; Johnson, C.E.; Morris, J.; Martin, G.; Lad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: mRNA
A; Residues: 1-107 <HAS>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            macrophage inflammatory protein 2 alpha precursor - human N; Alternate names: gro-beta; growth regulated protein beta; melanoma growth-stimulato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dр
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                                                                                                                                                                                                                                                                                                                                    A; Status: not compared with conceptual translation
                                                                                                                                                                                                                                                                                                                                                             A; Accession: A60407
                                                                                                                                                                                                                                                                                                                                                                                                                               A; Title: Monocyte adherence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Accession: A38290
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Title: Cloning and characterization of cDNAs for murine macrophage inflammatory pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C; Species: Homo sapiens (man)
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                                                                                                                                                                                                                                                                                                                                                                                               A; Reference number: A60407; MUID: 90257367
Query Match
Best Local Similarity
Matches 32; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   63
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   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 D.F.;
                               21.9%;
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   18;
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                      Score 128.5; DB 2;
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Pred. No. 1.2e-06;
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      Mismatches
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                                                              DB 2;
      35;
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      Indels
                                                              Length
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      11;
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   Gaps
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APPVSMRLLAAALLLLLLLALYTARVDGS-----KCKCSRKGPKIRYSDVKKLEMKPKYP 61

AAPSNPRLLRVALLLLLLVAASRRAAGAPLATELRCQCLQTLQGIHLKNIQSVKVKSPGP 67

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A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: mRNA A;Residues: 1-100 <RES> A;Cross-references: GB:S77604; NID:g998406; PIDN:AAB33749.1; PID:g998407 C;Superfamily: beta-thromboglobulin
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A;Stocession: I55614
A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       macrophage inflammatory protein-2 - rat
C;Species: Rattus sp. (rat)
C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 20-Aug-1999
C;Accession: I55614
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R; Feng, L.; Xia, Y.; Yoshimura, T.; Wilson, C.B.
J. Clin. Invest. 95, 1009-1017, 1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Reference number: A48988; MUID:93228656
A;Accession: D48988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: EMBL:X65647; NID:g56665; PIDN:CAA46599.1; PID:g56666 R;Nakagawa, H.; Ikesue, A.; Hatakeyama, S.; Kato, H.; Gotoda, T.; Komorita, N.; Watanabe
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A; Residues: 1-100 < DRI>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Title: Production of an interleukin-8-like chemokine by cytokine-stimulated rat NRK-49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R; Driscoll,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        macrophage inflammatory protein 2 -
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                                                                                                                                                                   Matches
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Best Local Similarity
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                                            N
                                                                                                8 APPVSMRLLAAALLLLLLALY---TARVDGS--KCKCSRKGPKIRYSDVKKLEMKPKYPH 62
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                                      APPTRQLLNAVLVLLLLLATNHQGTGVVVASELRCQCLTTLPRVDFKNIQSLTVTPPGPH 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPTRQLLNAVLVLLLLLATNHQGTGVVVASELRCQCLTTLPRVDFKNIQSLTVTPPGPH 61
                                                                                                                                                                                                Similarity
                                                                                                                                                                Conservative
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                                                                                                                                                                                         21.1%; Score 124; DB 2; 33.7%; Pred. No. 6.3e-06;
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                                                                                                                                                                   16;
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A; Molecule type: protein
A; Residues: 35-41, 'x', 43-49, 'x', 51-52, 'xx', 55-57 <GOL>
R; Schroeder, J.M.; Persoon, N.L.M.; Christophers, E.
J. Exp. Med. 1/1, 1091-1100, 1990
A; Title: Lipophlysaccharide-stimulated human monocytes secrete, apart from neutrophil ntity with melanoma growth stimulatory activity.
A; Reference number: A47626; MUID: 90217938
A; Accession: A47626
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Biochem. J. 259, 585-588, 1989
A;Title: Inflammatory cytokines induce synthesis and secretion of gro protein and a n A;Reference number: S03975; MUID:89246368
A;Accession: S03976
                                                                                                                                                   A; Molecule type: protein A; Residues: 35-62 < PRO>
                                                                                                                                                                                                         A;Reference number: A46519; MUID:93139489
A;Accession: B46519
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C;Date: 30-Jun-1989 #sequence revision 30-Jun-1989 #text_change 20-Aug-1999
C;Accession: $13669; A28414; $00983; B60401; $03976; A47626; B46519
R;Baker, N.E.; Kucera, G.; Richmond, A.
Nucleic Acids Res. 18, 6453, 1990
Nucleic Acids Res. 18, 6453, 1990
   A; Map position
                                A;Cross-references: GDB:120181; OMIM:155730
                                                            A; Gene: GDB:GRØ1
                                                                                            A; Experimental source: MG-63 C; Genetics:
                                                                                                                                                                                                                                                                J. Immunol. 15b, 1000-1010, 1993
A:Title: Identification of a novel granulocyte chemotactic protein (GCP-2) from human
                                                                                                                                                                                                                                                                                                                                      R; Proost, P.;
                                                                                                                                                                                                                                                                                                                                                                A; Experimental
                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: protein
A; Residues: 35-63, 'X', 65 < SC2>
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Biochem. J. 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Experimental source: dermal fibroblasts
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A; Residues: 35-42, 'X', 44, 'X', 46-48 <SCH>
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R;Schroeder, J.M.; Sticherling, M.; Hennelcke, H.H.; Preissner, W.C.; Christophers, V.Schroeder, J.M.; Sticherling, M.; Hennelcke, H.H.; Preissner, W.C.; Christophers, J. Immunol. 144, 223-2232, 1990
A;Title: IL-1alpha or tumor necrosis factor-alpha stimulate release of three NAP-1/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-107 < RIC>
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A;Accession: S00983
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R; Richmond, A.; Balentien, EMBO J. 7, 2025-2033, 1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-107 < ANI>
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A; Residues: 1-107 < BAK>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Status: preliminary
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9, 585-588, 1989
4q21-4q21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             necrosis factor-alpha stimulate release of three NAP-1/IL
                                                                                                                         osteosarcoma cells
                                                                                                                                                                                                                                                                                                                             C.; Conings,
                                                                                                                                                                                                                                                                                                                                Opdenakker, G.; Billiau, A.; Van Damm
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in transformed Chines

Spiess, J.; Bo

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NyAlternate names: growth-regulated protein gamma; macrophage inflammatory protein 2 bet C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 31-May-1991 #sequence_revision 27-Oct-1995 #text_change 20-Aug-1999
C;Accession: JH0282; B38290; C46519
R;Tekamp-Olson, P; Gallegos, C; Bauer, D.; McClain, J.; Sherry, B.; Fabre, M.; van Dev J. Exp. Med. 172, 911-919, 1990
A;Title: Cloning and characterization of cDNAs for murine macrophage inflammatory protein A;Reference number: JH0280; MUID:90354792
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F;1-34/Domain: signal sequence #status predicted <SIG>
F;35-107/Product: melanoma growth-stimulatory activity #status experimental <MAT>
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A; Residues: 1-26, 'G', 29-107 <HAS>
A; Residues: 1-26, 'G', 29-107 <HAS>
A; Cross-references: GB:M36821; NID:g183632; PIDN:AAA63184.1; PID:g183633
A; Cross-references: GB:M36821; NID:g183632; PIDN:AAA63184.1; PID:g183633
A; Proost, P; De Wolf-Peeters, C.; Conings, R.; Opdenakker, G.; Billiau,
J. Immunol. 150, 1000-1010, 1993
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F;1-34/Domain: signal sequence #status predicted <SIG>
F;35-107/Product: GRO-gamma #status experimental <MAT>
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gro-alpha precursor - mouse
N;Alternate names: gro protein;
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                                                                                                                                                                                                                                                                                                                                                                           Qγ
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                                                                    A32954
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                                                                                                                                                                                                 HCAQTEVIATIKN-----GKKACLNPASPMVQKIIE 98
                                                                                                                                                                                                                                                            HCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
                                                                                                                                                                                                                                                                                                                                                                                     APPVSMRLLAAALLLLLLLALYTARVDGS-----KCKCSRKGPKIRYSDVKKLEMKPKYP 61
                                                                                                                                                                                                                                                                                                                       AAPSNPRILRVALLILLIVAASRRAAGASVVTELRCQCLQTLQGIHLKNIQSVNVRSPGP 67
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   growth regulated protein; melanoma
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         growth-stimulating
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A;Cross references: GB:J04596; NID:g201042; PIDN:AAA40131.1; PID:g201043 R:Ryseck, R.P.; MacDonald-Bravo, H.; Mattei, M.G.; Bravo, R. Exp. Cell Res. 180, 266-275, 1989 A;Title: Cloning and sequence of a secretory protein induced by growth fa A;Reference number: JH0081; MUID:89078502 A;Accession: JH0081
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C;Date: 20-Oct-1989 #sequence_revision 20-Oct-1989 #text_change 20-Aug-1999
C;Accession: A32954; JH0081
R;Oquendo, P.; Alberta, J.; Wen, D.; Graycar, J.L.; Dervnck, R.: Stiles. C [
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Goodman, R.B.; Foster, D.C.; Mathewes, S.L.; Osborn, S.G.; Kuijper, J.L.; Forstrom, Biochemistry 31, 10483-10490, 1992
A;Title: Molecular cloning of porcine alveolar macrophage-derived neutrophil chemotac A;Reference number: A44253; MUID:93041741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     alveolar macrophage chemotactic factor-II (AMCF-II) intercrine-alpha protein C;Species: Sus scrofa domestica (domestic pig) C;Date: 30-Apr-1993 #sequence_revision 18-Nov-1994 #text_change 20-Aug-1999 C;Accession: B44253
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A;Title: The platelet-derived growth factor-inducible KC gene encodes a secretory pro
                                                                                                                                                                                                                                                                                                                                                                                                    A; Experimental source: alveolar macrophage A; Note: sequence extracted from NCBI backbone (NCBIN:117417, NCBIP:117418) C; Superfamily: beta-thromboglobulin
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61
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IHPKMISDLQVIPAGPQCSKAEVIATLKN-----GKEVCLDPKAPLIKKIVQ 107
                                                             IRYSDVKKLEMKPKYPHCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
                                                                                                                          MRLLTSRATRVPSPSGLLCAVLAMLLLTPSGPLASASPIEAAEAAVVRELRCMCLTTTPG 60
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Pred. No. 0.00014;
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                                                                                                                                                                                                                                                                       C;Superfamily: beta-thromboglobulin
C;Keywords: cytckine; disulfide bond
F;1-24/Domain: signal sequence #status predicted <SIG>
F;25-96/Product: neutrophil chemo-attractant Gro prote
                                                                                                                                                                                                                                                                                                                                                                                        A;Gene: gro; KC
A;Introns: 24/1; 65/2; 92/2
                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            C:Comment: This protein has chemotactic activity for neutrophils and has melanoma growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Note: sequence extracted from NCBI backbone (NCBIP:129131) R:Hanzawa, H.; Haruyama, H.; Watanabe, K.; Tsurufuji, S. FEBS Lett. 354, 207-212, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R; Watanabe, K.; Konishi, K.; Fujioka, M.; Kinoshita, S.; Nakagawa, H.
J. Biol. Chem. 264, 19559-19563, 1989
A;Title: The neutrophil chemoattractant produced by the rat kidney epithelioid A;Reference number: A34481; MUID:90062049
A;Accession: A34481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: protein A; Residues: 25-96 <HAN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Title: The three dimensional structure of rat cytokine CINC/Gro in solution by homonuous, Reference number: S51214; MUID: 95046335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: protein A; Residues: 25-57 <NA2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Residues: 25-96 <WAT>
R;Nakagawa, H.; Ikesue, A.; Hatakeyama, S.; Kato, H.; Gotoda, T.; Komorita, N.; Watanabo Biochem. Pharmacol. 45, 1425-1430, 1993
A;Title: Production of an interleukin-8-like chemokine by cytokine-stimulated rat NRK-49A;Reference number: A48988; MUID:93228656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: mRNA
A; Residues: 1-32,'S',34-96 <HUA>
A; Cross-references: GB: M86536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Konishi, k.; ..... 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Accession: S51214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Contents: annotation; conformation by (1) H-NMR, residues 25-96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Experimental source: kidney, NRK-49F fibroblasts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Accession: B48988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: protein A; Residues: 25-57 < NAK>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Note: the authors translated the codon AGT for residue 33 as Cys, AAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Title: Rat KC cDNA cloning and mRNA expression in lung macrophages and fibroblasts.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Huang, S.; Paulauskis, J.D.; Kobzik, L.
Biochem. Biophys. Res. Commun. 184, 922-929, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: DDBJ:D11445; NID:g391854; PIDN:BAA02009.1; PID:g220755
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 20-Jun-2000 C; Accession: JN0572; JQ1519; A34481; A48988; B48988; S51214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               neutrophil chemo-attractant Gro protein precursor - rat
N;Alternate names: CINC; cytokine-induced neutrophil chemoattractant; interleukin-8-like
C;Species: Rattus norvegicus (Norway rat)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Note:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Experimental source: kidney, NRK-49F fibroblasts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Experimental source: alveolar macrophage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Reference number: JQ1519; MUID:92246987
A; Accession: JQ1519
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Accession: JN0572
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                                                                                                                                                                                                Query Match
                                                                      15 LLAAALLLLLALYTAR-----VDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKM 67
                                                                                                                                                                      Local
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                    LLCAALPVLA----TSRQATGAPVANELRCQCLQTVAGIHFKNIQSLKVMPPGPHCTQTE 63
                                                                                                                                                                      Similarity
                                                                                                                                      Conservative
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                                                                                                                                                             16.5%;
27.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              from NCBI backbone (NCBIP:129132)
                                                                                                                                      20;
                                                                                                                                                             Score 97;
Pred. No.
                                                                                                                                                             0.004;
                                                                                                                                                                                        DB 2; Length 96
                                                                                                                                                                                                                                                                    protein #status experimental <CYT>
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C;Species: Rattus sp. (rat)
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 16-Feb-1997
                                                             macrophage inflammatory protein-2 - rat (fragment)
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C;Superfamily: beta-
C;Keywords: cytokine
                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: mRNA
A; Residues: 1-125 <FAR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biochem. Biophys. Res. Commun. 192, 223-230, 1993
A;Title: HuMig: a new human member of the chemokine family of cytokines
A;Reference number: JN0470; MUID:93236577
                                                                                                                                                                                                                                                                                                                                                                       A;Cross_references: GB:X72755; GB:S60728; NID:g311375; PIDN:CAA51284.1; PID:g311376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    interferon gamma-induced protein HuMIG - human C:Species: Homo sapiens (man)
                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Accession: JN0470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R; Farber, J.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 28-May-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Proc. Natl. Abad. Sci. U.S.A. 87, 5238-5242, 1990
A;Title: A matrophage mRNA selectively induced by gamma-interferon encodes a member o
A;Reference number: A35766; MUID:90319087
A;Accession: A35766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    platelet factor 4, interferon gamma-induced, precursor - mouse
C:Species: Mus musculus (house mouse)
C:Date: 05-Oct-1990 #sequence_revision 05-Oct-1990 #text_change 20-Aug-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross_references: GB:M34815; NID:g199692; PIDN:AAA39706.1; PID:g199693
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A; Residues: 1-126 <FAR>
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Best Local
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                                               74 SVSHYRGQEHCLHPKLQSTKRFI-KWYNAWNEKRR 107
                                                                                                                                             15 LLAAALLLLLLALYTARVDGSKCKC-SRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITTK 73
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                                                                                                 9 LLGTILLVLIGVQGTPVVRKGRCSCISTNQGTIHLQSLKDLKQFAPSPSCEKIEIIATLK 68
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--GVQTCLNPDSADVKELIKKWEKQVSQKKK 98
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                                                                                                                                                                                                                          14.9%; Score 87.5; DB 29.5%; Pred. No. 0.05;
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C;Accession: I51886
R;Huang, S.; Pathol. 141, 981-988, 1992
A;Title: Expression of macrophage inflammatory protein-2 and KC mRNA in pulmonary inflam A;Reference number: I51886; MUID:93035653
A;Accession: I51886; MUID:93035653
A;Accession: I51886; MUID:93035653
A;Accession: I51886; MUID:93035653
A;Accession: I51886; MUID:93035653
A;Reference number: I51886; MUID:93035653
A;Residues: Prediminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-53 <RES>
A;Cross-references: GB:S4585; NID:9257054
C;Superfamily: beta-thromboglobulin
C;Superfamily: beta-thromboglobulin
Ouery Match
Best Local Similarity 29.6%; pred. No. 0.035;
Matches 16; Conservative 13; Mismatches 20; Indels 5; Gaps 1;
Matches 16; Conservative 13; Mismatches 20; Indels 5; Gaps 1;
Db PRVDFKNIQSLTVTPPGPHCAQTEVIATLKD-----GHEVCLNPEAPLVQRIVQ 53
Search completed: January 29, 2002, 16:04:46
Job time: 72 sec
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GenCore version 4.5

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OM protein - protein search, using sw model

Run on:

January 29, 2002, 16:03:34; Search time 11.67 Seconds (without alignments)
348.740 Million cell updates/sec

Title:
Perfect score: 587
Sequence: 1 MSLLPRRAPPVSMRLLAAAL.....TKRFIKWYNAWNEKRRVYEE 111

Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5

Searched: 100059 seqs, 36664827 residues

Total number of hits satisfying chosen parameters: 100059

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Database · Swissprot 30.*

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database : SwissProt_39:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Score 527 489 143.5 138 138 139 143.5 117.5 117.5 1117.5
Watch I 89.8 3 3 89.3 3 221.7 221.7 221.7 120.0 20.0 20.0 20.0 20.0 20.0 20.0 20.
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PP9967X4F118886331
Description 095715 homo 09wq5 mus 010747 ratt p10889 mus p109340 cric p109348 ratt p10875 bos p46675 bos p46676 bos p19876 homo 046676 bos p19876 homo p19876 ratt 055235 cavi p12850 mus p14095 ratt p18340 mus p10746 rats p
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PRINTS; PR00436; INTERLEUKIN8.
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Biochem. Biophys. Res. Commun. 255:703-706(1999)
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                                                                                                                   PRINTS; PR00436; INTERLEUKIN8. PRINTS; PR00437; SMALLCYTKCXC.
                                                                                                                                                                                                                                                                                                                                                               use by non-profit institutions as long modified and this statement is not removed
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01-OCT-1996 (Rel. 34, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
MACROPHAGE INFLAMMATORY PROTEIN-2-BETA PRECURSOR (MIP2-BETA) (CINC-
                                                                                                                                                                                                                   InterPro; IPR001811; Chemokine_IL8.
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"Identification of cytokine-induced neutrophil chemoattractants
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91.9%;
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MACROPHAGE INFLAMMATORY PROTEIN-
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                    Shao W., Jerva L.F., West J., Lolis E., Schweitzer B.I.;
"Solution structure of murine macrophage inflammatory protein-2.";
Biochemistry 37:8303-8313(1998).
-!- PUNCTION: CHEMOTACTIC FOR HUMAN POLYMORPHONUCLEAR LEUKOCYTES BUDES NOT INDUCE CHEMOKINESIS OR AN OXIDATIVE BURST.
-!- SUBUNT: HOWOTETRAMER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Mammalia; Eutheria;
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01-JUL-1989
                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
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                                                                                                       EMBL; X53798; CAA37807.1; PIR; JH0200; JH0200.
                                                                                                                                                                                   entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                      use by non-profit institutions as long modified and this statement is not removed.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Proc. Natl. Acad. Sci. U.S.A. 86:612-616(1989).
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erPro; IPR001811; Chem
erPro; IPR001089; Smal
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p09340;
01-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last sequence
20-AUG-2001 (Rel. 40, Last annotati
20-AUG-201 (Rel. 40, Last annotati
20-MTH REGULATED PROTEIN PRECURSOR.
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Eukaryota; Metazoa; Chordata; Crania
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PRINTS; PR00437; SMAL
           SIGNAL
CHAIN
                                                PRINTS; PRO0436; INTERLEUKIN8.
PRINTS; PRO0437; SMALLCYTKCXC.
SMART; SM00199; SCY; 1.
PROSITE; PS00471; SMALL_CYTOKI
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InterPro; IPR001089; Small_cytokine_CXC
Pfam; PF00048; IL8; 1.
                                                                                                                                                                                                                                    the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                   Anisowick A., Bardwell L., Sager R.;

"Constitutive overexpression of a growth-regulated gene
"transformed Chinese hamster and human cells.";

Proc. Natl. Acad. Sci. U.S.A. 84:7188-7192(1987).

-i- FUNCTION: HAS CHEMOTACTIC ACTIVITY FOR NEUTROPHILS.
                                                                                                                                                    PIR; B28414; B28414.
HSSP; P10889; 1MI2.
                                                                                                                                                                                                         entities requires a license agreement (son send an email to license@isb-sib.ch)
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Mammalia¦ Eutheria; Rodentia; Sciurognathi; Muridae; Cricetinae;
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DISULFID
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PRO0437; SMALLCYTKCXC.
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                                                   SMALL_CYTOKINES_CXC; 1.
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Last annotation updat
                         Inflammatory response; POTENTIAL.
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Pred. No. 5.6e-08;
GROWTH REGULATED PROTEIN BY SIMILARITY.
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                                                                              "Nuclear magnetic resonance solution structure of truncated human GRObeta [5-73] and its structural comparison with CXC chemokine family members GROalpha and IL-8.";
J. MOI. Biol. 294:1065-1072(1999).
-i- FUNCTION: PRODUCED BY ACTIVATED MONOCYTES AND NEUTROPHILS AND EXPRESSED AT SITES OF INFLAMMATION.
-i- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUR-Histiocytic lymphoma;
MEDLINE=90354792; PubMed=2201751;
MEDLINE=90354792; PubMed=2201751;
Tekamp-Olson P., Gallegos C., Bauer D., McClain J., Sherry B.
Fabre M., van Deventer S., Cerami A.;
"Cloning and characterization of cDNAs for murine macrophage inflammatory protein 2 and its human homologues.";
J. Exp. Med. 172:911-919(1990).
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P19875; Q9UBB8;
01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
20-AUG-2001 (Rel. 40, Last annotation update)
MACROPHAGE INFLAMMATORY PROTEIN-2-ALPHA PRECURSOR (MIP2-ALPHA) (GROWTH
                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE OF 35-107 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Haskill S., Peace A., Morris J., Sporn S.A., Anisowicz A., Lee S.W. Smith T., Martin G., Ralph P., Sager R.; "Identification of three related human GRO genes encoding cytokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Cloning and sequencing of a new gro transcript from activated human monocytes: expression in leukocytes and wound tissue.";
Mol. Cell. Biol. 10:5596-5599(1990).
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                                                                                                                                                                                                                                                                                            Qian Y.Q., Johanson K.O., McDevitt P.;
                                                                                                                                                                                                                                                                                                                          MEDLINE=20069929; PubMed=10600366;
                                                                                                                                                                                                                                                                                                                                                      STRUCTURE BY NMR OF 39-107
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Eukaryota; Metazoa;
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                                                            C-X-C) (CHEMOKINE CXC).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Natl. Acad. Sci. U.S.A. 87:7732-7736(1990).
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Pred. No. 3.1e-07;
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Best Local
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                                                                                                                                                                                                                                            Yoshimura T., Modi W.S.;
"Isolation of novel GRO genes, and a phylogenetic analysis chemokine subfamily in mammals.";
Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases.
-!- EUNCTION: HAS CHEMOTACTIC ACTIVITY FOR NEUTROPHILS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GRO_SHEEP
O46678;
                         the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for con
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
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Bovidae; Caprinae; Ovis.
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30-MAY-2000 (Rel.
20-AUG-2001 (Rel.
                                                                                                                                                                                                                     -!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE
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the European Bioinformatics Institute. There are no restrictions
use by non-profit institutions as long as its content is in
modified and this statement is not removed. Usage by and for com
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InterPro; IPR001089; Small_cytokine_CXC
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107 AA;
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01-APR-1993 (Rel. 25, Last sequence update)
10-AUG-2001 (Rel. 40, Last annotation update)
MACROPHAGE INFLAMMATORY PROTEIN 2 PRECURSOR (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRINTS; PRO0436; INTERLEUKIN8. PRINTS; PRO0437; SMALLCYTKCXC. SMART; SM00199; SCY; 1.
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                                                                                                                      STRAIN=CD; TISSUE=Lung; Farone A., Farone M., Sh Submitted (MAR-1996) to
                                                                                                                                                                                  MEDLINE=95189993; PubMed=7883948; Feng L., Xia Y., Yoshimura T., Wilson C.B.; "Modulation of neutrophil influx in glomerulonephritis in the rat with anti-macrophage inflammatory protein-2 (MIP-2) antibody."; J. Clin. Invest. 95:1009-1017(1995).
                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                          Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                    Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chor
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InterPro; IPR001089; Small_cytokine
                                                                TISSUE=Lung;
MEDLINE=93035653; PubMed=1415488;
                                                                                                                                                                                                                                               STRAIN-FISCHER;
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                                                                                                                                                                                                                                                                                                          STRAIN=FISCHER 344; TISSUE=Lung;
                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=10116;
SEQUENCE OF 32-100
                        pulmonary inflammation.";
Am. J. Pathol. 141:981-988(1992)
                                                Huang S., Paulauskis J.D., Godleski J.J., Kobzik "Expression of macrophage inflammatory protein-2
                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                   Submitted (APR-1992)
                                                                                                SEQUENCE OF 39-91 FROM N.A.
                                                                                                                                                            SEQUENCE FROM N.A.
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                                                              Godleski J.J., Kobzik L.;
                                                                                                                         EMBL/GenBank/DDBJ
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BY SIMILARITY.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nakagawa H., Shiota S., Takano K., Shibata F., Kato H.;
"Cytokine-induced neutrophil chemoattractant (CINC)-2 alpha, a no
"Cytokine-induced neutrophil chemoattractant (CINC)-2 alpha, a no
member of rat GRO/CINCS, is a predominant chemokine produced by
lipopolysaccharide-stimulated rat macrophages in culture.";
Biochem. Biophys. Res. Commun. 220:945-948(1996).
Biochem. Biophys. Res. Commun. 220:945-948(1996).

FUNCTION: CHEMOTACTIC FOR HUMAN POLYMORPHIONUCLEAR LEUKOCYTES
DOES NOT INDUCE CHEMOKINESIS OR AN OXIDATIVE BURST. CONTRIBUT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Identification of cytokine-induced neutrophil chemoattractants (CINC), rat GRO/CINC-2 alpha and CINC-2 beta, produced by granulation tissue in culture: purification, complete amino acid sequences and characterization.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=94318061; PubMed=8043001;
Nakagawa H., Komorita N., Shibata
Fujioka M., Kato H.;
                                                                                                                                                                                                                                                                                                 CHAIN
DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biochem.
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SMART; SM00199; SCY; 1.
PROSITE; PS00471; SMALL_CYTOKINES_CXC; 1.
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HSSP; P10889; 1MI2.
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                                                                62 CAQTEVIATLKD-----GHEVCLNPEAPLVQRIVQ
                                                                                            63 CEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SUBUNIT: HOMOTETRAMER.
TISSUE SPECIFICITY: AT LEAST EXPRESSED IN THE LUNG AND TRACHEA.
INDUCTION: IN LUNG, BY LIPOPOLYSACCHARIDE OR INFLAMMATION.
SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C-X-¢) (CHEMOKINE CXC).
                                                                                                                                                       APPVSMRLLAAALLLLLLALY---TARVDGS--KCKCSRKGPKIRYSDVKKLEMKPKYPH 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SWISS-PROT entry is copyright. It is produced through the Swiss Institute of Bioinformatics and the EN
                                                                                                                           APP|TRQLLNAVLVLLLLLATNHQGTGVVVASELRCQCLTTLPRVDFKNIQSLTVTPPGPH
                                                                                                                                                                                                                                                                                                                                                                                                                          PF00048; IL8; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   an
                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                         PRO0436; INTERLEUKIN8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and this statement is not removed. Usage by and for commercial requires a license agreement (See http://www.isb-sib.ch/announce/an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ROPHIL ACTIVATION DURING INFLAMMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               J. 301:545-550(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STAR;
                                                                                                                                                                                                                                                                                                                                                                                                                                       IPR001811; Chemokine_IL8.
IPR001089; Small_cytokine_CXC
                                                                                                                                                                                                                                                                                                                                              Chemotaxis; Inflammatory response; Signal.
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                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                     3852
                                                                                                                                                                                                                                                                   AA;
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                                                                                                                                                                                                                                                                     10783
                                                                                                                                                                                                         21.1%;
                                                                                                                                                                                                                                                                     MW;
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                                                                                                                                                                                                       Score 124;
Pred. No. 1
                                                                                                                                                                                                                                                                                    BY SIMILARITY
BY SIMILARITY
                                                                                                                                                                                                                                                                                                                  MACROPHAGE INFLAMMATORY PROTEIN 2.
                                                                                                                                                                                                                                                                   4299DFE68FF37A9B CRC64
                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                         DB 1;
.7e-06;
                                                                  91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 as its content
                                                                                                                                                                                                                    Length 100;
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Cytokine; Growth factor; I
SIGNAL
1 29
CHAIN 30
DISULFID 39
DISULFID 39
DISULFID 41
81
Submitted (JUL-1994) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: PLAYS A ROLE IN MONOCYTE ADHESION TO THE ENDOTHELIUM.
-!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE C-X-C) (CHEMOKINE CXC).
                                                                                                                                                               Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                 Ol-FEB-1996 (Rel. 33, Created)
Ol-FEB-1996 (Rel. 33, Last sequence update)
Ol-FEB-1996 (Rel. 33, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DISULFID
DISULFID
SEQUENCE
                                                                                   Schwartz D., Chaverri-Alamada L., Be Quisomoro D., Fang J., Tekamp-Olson
                                                                                                                                                                                                                OI-FEB-1996 (Rel. 33, Last annotation update)
GROWTH REGULATED PROTEIN HOMOLOG PRECURSOR (GRO HOMOLOG).
                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                   01-FEB-1996
                                                                                                                                                                                                                                                                                     P47854;
                                                                                                                                                                                                                                                                                                                      RABIT
                                                                     Territo M.;
                                                                                 Quisomoro D.,
                                                                                                                                                   NCBI_TaxID=9986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
GROWIH REGULATED PROTEIN HOMOLOG GAMMA PRECURSOR (GRO-GAMMA).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRINTS;
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Pfam; PF00048; IL8; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bovidae; Bovinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; U95811; AAB93927.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bos taurus (Bovine).
                                                                                                                                                                                                                                                                                                                                                                                                              60 YPHCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFI 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                                                                                                   RABIT
                                                                                                                                                                                                                                                                                                                                                                                 GPHCDQTEVIATLKT-----GQEVCLNPAAPMVKKII 93
                                                                                                                                                                                                                                                                                                                                                                                                                                              APAASSAPRILRAAMILLILVAAGRRAAGAPVVNELRCQCLQTLQGIHLKNIQSVKVTTP 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPVSM -- RLLAAALLLLLLALYTARVDGS ----- KCKCSRKGPKIRYSDVKKLEMKPK 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; PR00437; SMALLCYTKCXC SM00199; SCY; 1.
E; PS00471; SMALL_CYTOK:
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                                                                                                                                                                                                                                                                                                   STANDARD;
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factor; Inflammatory response; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10393 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 122.5; Db _,
Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GROWTH REGULATED PROTEIN HOMOLOG GAMMA. BY SIMILARITY. BY SIMILARITY.
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                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            942CD6897C21EDE9 CRC64;
                                                                                                  Berliner J.,
                                                                                   P.,
                                                                                                                                                                                                                                                                                                 104 AA
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                                                                                   Lusis
                                                                                   J.,
                                                                                                Kirchgessner T.,
                                                                                 Fogelman
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GROB_BOVIN
IID GROB_BAC
AC 046677
DT 15-DEC
DC Eukary
OC Mammal
OC Bovida
OX NCEI_T
RN [1]
RP SEQUEN
RA YOShim
RL Submit
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Best Local
                                                                                                                                                                                                                                                                                                     use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
GROWITH REGULATED PROTEIN HOMOLOG BETA PRECURSOR (GRO-BETA).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DISULFID DISULFID
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SIGNAL
        PRINTS; PR00436; PRINTS; PR00437;
                                                                                 Pfam; PF00048; IL8;
                                                                                                                  InterPro; IPR001811; Chemokine_IL8.
InterPro; IPR001089; Small_cytokine
                                                                                                                                                                                                       EMBL; U95813; AAB93929.1; -
                                                                                                                                                                                                                                                                                or send an email to license@isb-sib.ch).
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PRINTS; PRO0437; SMALLCYTKCXC.
SMART; SM00199; SCY; 1.
PROSITE; PS00471; SMALL_CYTOKINES_CXC; 1.
CYTOKINE; Growth factor; Inflammatory response; Signal.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9913;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GROB_BOVIN O46677;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
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InterPro; IPR001089; Small_cytokine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              62
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C-X-C) (CHEMOKINE CXC).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPVSMRLLAAALLLLLLALYTARVDGS-----KCKCSRKGPKIRYSDVKKLEMKPKYP 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFI 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATAAAPRLLRAALLLLLLVAAGRRAAGAPVVNELRCQCLQTLQGIHLKNIQSVKVTTPGP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bovinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         104 AA;
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                                                INTERLEUKIN8
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66 B
82 B
; 10900 MW;
                                                                                                                      Small_cytokine_CXC
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Pred. No. 6.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BY SIMILARITY.
BY SIMILARITY.
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SMALLCYTKCXC

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RESULT 12
GRO_HUMAN
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Matches 30
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01-MAR-1989
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DISULFID
SEQUENCE
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01-MAR-1989 (Rel. 10, Last sequence update)
20-AUG-2001 (Rel. 40, Last annotation update)
GROWTH REGULATED PROTEIN PRECURSOR (MELANOMA GROWTH STIMULATORY
ACTIVITY) (MGSA) (NEUTROPHIL-ACTIVATING PROTEIN 3) (NAP-3).
SCYBI OR GRO1 OR GROA OR GRO OR MGSA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.

MEDLINE-88328991; PubMed=2970963;

MICHINE-88328991; PubMed=2970963;

MICHINE-88328991; PubMed=2970963;

MICHINE-88328991; PubMed=2970963;

MEDLINE-88328991; PubMed=2970963;

MEDLINE-8832891; PubMed=2970963;

MEDLINE-88328991; PubMed=2970963;

MEDLINE-8832891; PubMed=2970963;
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                                                                                                     Schroeder J.-M., Persoon N.L.M., Christophers E., "Lipopolysaccharide-stimulated human monocytes secrete, apart from neutrophil-activating peptide l/interleukin 8, a second neutrophil-activating protein. NH2-terminal amino acid sequence identity with
                                                                                                                                                                                                                                                                             "Nucleotide sequence of the human melanoma growth stimulatory activity (MGSA) gene."; Nucleic Acids Res. 18:6453-6453(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Constitutive overexpression of a growth-regulated transformed Chinese hamster and human cells."; Proc. Natl. Acad. Sci. U.S.A. 84:7188-7192(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
MEDLINE=88041072; PubMed=2890161;
Anisowicz A., Bardwell L., Sager
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                growth stimulatory activity, a growth factor structurally related
beta-thromboglobulin.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GRO_HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytokine;
MEDLINE=89246368;
                   SEQUENCE OF
                                                                                           melanoma
                                                                                                                                                                                                        MEDLINE=90217938; PubMed=2182761;
                                                                                                                                                                                                                                   SEQUENCE OF 35-65.
                                                                                                                                                                                                                                                                                                                                                   Baker N.E., Kucera G.,
                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=91057157; PubMed=2129556;
                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SMART; SM00199; SCY; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFI 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATAAAPRILRAAMLLLLLVAAGRRAAGAPVVNELRCQCLQTLQGIHLKNIQSVKVTTPGP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7:2025-2033(1988).
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                                                                growth stimulatory activity.";
Med. 171:1091-1100(1990).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Growth factor;
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104 AA;
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PubMed=2655583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SMALL_CYTOKINES_CXC;
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                                                                                                                                                                                                                                                                                                                                                     Richmond A.;
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40A8C06A64D67F7B CRC64;
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GROWTH REGULATED PROTEIN HOMOLOG BETA.
BY SIMILARITY.
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Query Match
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EMBL; X12510; CAA31027.1; -.
EMBL; X54489; CAA38361.1; -.
PIR; A28414; A28414
PIR; 80988; S00983.
PIR; S03976; S03976.
PIR; S13669; S13669.
                                                                DISULFID DISULFID SEQUENCE
                                                                                                                                                                                                        SMART; S
PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced between the Swiss Institute of Bioinformatics and between the Swiss Institute. There are
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MEDLINE=94376296; PubMed=8089846;
Fairbrother W.J., Reilly D., Colby T.,
"The solution structure of melanoma ground of the structure of melanoma ground of the structure of the solution of the structure of the solution structure of the structure of the solution of the solution of the structure of the solution of the
                                                                                                                                                                                                                                                                                                                                                                                         PDB;
PDB;
PDB;
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J. Biol. Chem. 269:32909-32915(1994).
-i- EUNCTION: HAS CHEMOTACTIC ACTIVITY FOR NEUTROPHILS. MAY PLAY A ROLE IN INFLAMMATION AND EXERTS ITS EFFECTS ON ENDOTHELIAL CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Inflammatory cytokines induce synthesis and secretion of gro protein and a newtrophil chemotactic factor but not beta 2-microglobulin in human sympovial cells and fibroblasts.";
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Wen D., Rowland A., Derynck R.;
                                                                                                                                        CHAIN
                                                                                                                                                        SIGNAL
                                                                                                                                                                                   Cytokine
                                                                                                                                                                                                                                                                                                 InterPro: IPR001089; Small_cytokine_CXC
Pfam; PF00048; IL8; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C-X-C) (CHEMOKINE CXC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                       1MGS; 30-SEP-94
1MSG; 31-MAR-95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E.E., Mason
                                                                                                                                                                                                        SM00199; SCY;
E; PS00471; SM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8:1761-1766(1989)
                                                                                                                                                                                                                                                                      PRO0436; INTERLEUKIN8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       requires a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IN INFLAMMATION AND EXERTS AUTOCRINE FASHION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         growth stimulating activity (MGSA) by NMR spectroscopy.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FUNCTION.
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                                                                                                                                                                                Growth factor; Inflammatory response; Signal;
                                                                                                                                                                                                                                                                                                                                                                                         ; 31-MAR-95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clark-Lewis I.,
                                                                                                                                                                                                                                                                                                                                              IPR001811; Chemokine_IL8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      330:302-306(1993).
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                                                                     107
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                                                                     AA;
                                                                                                                                                                                                                                                   SMALLCYTKCXC
                                                                                                                                                                                                        SMALL_CYTOKINES_CXC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PubMed=7806518;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PubMed=8397104;
                                                                     11301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     license agreement (See http://www.isb-sib.ch/announce/
    20.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of
                                                                     WW.
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    Score 117.5;
                                                                                                                                        GROWTH REGULATED PROTEIN
                                                                     17048A6B4D765CA2 CRC64;
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    DВ
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                                                                                                                                                                                     3D-structure.
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Best Local Similarity

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RESULT 13
MIZB_HUMAN
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AC P19876
DT 20-AUG
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                                                                                                                                                                                                                                InterPro; IPR001811; Chemokine_IL8
InterPro; IPR001089; Small_cytokine
Pfam; PF00048; IL8; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE-90354792; PubMed-2201751; Tekamp-Olson P., Gallegos C., Bauer D., McClain J., Sherry B. Fabre M., van Deventer S., Cerami A.; "Cloning and characterization of cDNAs for murine macrophage inflammatory protein 2 and its human homologues."; J. Exp. Med. 172:911-919(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
01-FEB-201 (Rel. 40, Last annotation update)
MACROPHAGE INFLAMMATORY PROTEIN-2-BETA PRECURSOR (MIP2-BETA) (GROWTH
                                                           SIGNAL
                                                                                                                                         PRINTS; PR00436; INTERLEUKIN8.
PRINTS; PR00437; SMALLCYTKCXC.
SMART; SM00199; SCY; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Haskill S., Peace A., Morris J., Sporn Smith T., Martin G., Ralph P., Sager R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Ve
Mammalia; Eutheria; Primates; Catarrhini;
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01-FEB-1991
                                                                                   Cytokine; Chemotaxis; Inflammatory response; Signal.
                                                                                                                    PROSITE;
                                                                                                                                                                                                                                                                                                                          MIM; 139111;
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PIR; JH0282; JH0282.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TISSUE-Histiocytic lymphoma;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=91017578;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Identification of three related human GRO genes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C. Natl. Acad. Sci. U.S.A. 87:7732-7736(1990).
FUNCTION: MAY PLAY A ROLE IN INFLAMMATION AND EXERT ITS EFFECTS
ON ENDOTHELIAL CELLS IN AN AUTOCRINE FASHION.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            M36821; AAA63184.1;
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                                                                                                                 PS00471; SMALL_CYTOKINES_CXC; 1.
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107
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  MACROPHAGE INFLAMMATORY PROTEIN-2-BETA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                                                                                                                                                         PRINTS; PRO0436; INTERLEUKIN8.
PRINTS; PRO0437; SMALLCYTKCXC.
SMART; SM00199; SCY; 1.
PROSITE; PS00471; SMALL_CYTOKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-DEC-1998 (Rel. 37, Last annotation update) GROWIH REGULATED PROTEIN HOMOLOG ALPHA PRECURSOR (GRO-ALPHA).
                                                                                                                                                                                                                                                                                                                                                                     CHAIN
                                                                                                                                                                                                                                                                                                                                                                                            SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                     Cytokine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pfam; PF00048; IL8; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR001811; Chemokine_IL8.
InterPro; IPR001089; Small_cytokine_CXC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; U95812; AAB93928.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bos taurus (Bovine).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-DEC-1998 (Rel.
15-DEC-1998 (Rel.
15-DEC-1998 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     046676;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GROA_BOVIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yoshimura T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bovidae; Bovinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mammalia; Eutheria;
                           62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    68 HCAQTEVIATIKN-----GKKACLNPASPMVQKIIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            62 HCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8 AAPSNPRLLRVALLLLLLVAASRRAAGASVVTELRCQCLQTLQGIHLKNIQSVNVRSPGP 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8 APPVSMRLLAAALLLLLLALYTARVDGS-----KCKCSRKGPKIRYSDVKKLEMKPKYP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C-X-C) (CHEMOKINE CXC).
HCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFI 96
                                                                           ATAAAPRLLRAAMLFLLLVAAGRRAAGAPVVNELRCQCLQTLQGIHLKNIQSVKVTTPGP 64
                                                                                                                          APPVSMRLLAAALLLLLLALYTARVDGS------KCKCSRKGPKIRYSDVKKLEMKPKYP 61
                                                                                                                                                                          29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29;
                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 30.:
29; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                  Growth factor; Inflammatory response; Signal.
                                                                                                                                                                                                                                                                                               104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (Rel. 37, Created)
(Rel. 37, Last sequence update)
(Rel. 37, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                107 AA;
                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                      31
40
42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Modi W.S.;
                                                                                                                                                                                                                                                                                               AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bos.
                                                                                                                                                                                                                                                                                             104 c
66 E
82 E
10984 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                      SMALL_CYTOKINES_CXC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           69 B
85 B
28 A
11342 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chordata; Craniata; Vertebrata; Euteleostomi; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19.8%;
30.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
                                                                                                                                                                                                Score 113.5;
Pred. No. 2.
                                                                                                                                                                                                                                                                                             GROWTH REGULATED PROTEIN HOMOLOG ALPHA.
BY SIMILARITY.
BY SIMILARITY.
1002CAC064DB1F76 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 116.5;
Pred. No. 1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A
                                                                                                                                                                                                                                                                                                                                                                                              POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97A69946B7F1F070 CRC64;
                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              104
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.1e-05
                                                                                                                                                                                                .3e-05;
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                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                          37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pecora; Bovoidea;
                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 107;
                                                                                                                                                                          Indels
                                                                                                                                                                                                                       104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11;
                                                                                                                                                                          11;
                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61
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B

65 HCDQTEVIASLKT-----GQEVCLNPTAPMVKKII 94

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GRO_MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT
                             EMBL; J04596; AAA40131.1; -.
EMBL; U20634; AAB03376.1; JOINED.
EMBL; U20627; AAB03376.1; JOINED.
EMBL; S79767; -: NOT_ANNOTATED_CDS.
PIR; J32954; A32954.
PIR; J40081; J40081.
HSSP; P09341; IMSH.
MGD; MGI:108068; GrOl.
InterPro; IPR001811; Chemokine_IL8.
InterPro; IPR001811; Chemokine_IL8.
InterPro; IPR00189; Small_cytokine_CXC.
Pfam; PF00048; IL8; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P12850;
01-OCT-1989
01-OCT-1989
20-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                     "Two structurally distinct kappa B sequence motifs cooperatively control LPS-induced KC gene transcription in mouse macrophages.";

J. Immunol. 155:3593-3600(1995).

-!- FUNCTION: HAS CHEMOTACTIC ACTIVITY FOR NEUTROPHILS. CONTRIBUTES TO NEUTROPHIL ACTIVATION DURING INFLAMMATION (BY SIMILARITY).

-!- INDUCTION: BY PLATELET DERIVED GROWTH FACTOR. IN LUNG, BY LIPOPOLYSACCHARIDE OR INFLAMMATION (BY SIMILARITY).

-!- SIMILARITY: BELONGS TO THE INTERCRINE ALPHA FAMILY (SMALL CYTOKINE C.Y.C.) (CHEMOKINE CXC).
                                                                                                                                                                                                                                                       use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-OCT-1989 (Rel. 12, Created)
01-OCT-1989 (Rel. 12, Last sequence update)
20-AUG-2001 (Rel. 40, Last anotation update)
GROWTH REGULATED PROTEIN PRECURSOR (PLATELET-DERIVED GROWTH FACTOR-INDUCIBLE PROTEIN KC) (SECRETORY PROTEIN N51).
                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bozic C.R., Kolakowski L.F. Jr., von Uexkull C., Garcia-Rodriguez M., Conklyn M.J., Breslow R., Showell H.J., Gerard N.P., Gerard C.; Submitted (FEB-1995) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ryseck R.P., Macdonald-Bravo H., Mattei M.-G., Bravo R.; "Cloning and sequence of a secretory protein induced by growth factors in mouse fibroblasts.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A. MEDLINE=89078502; Pubmed=2909392;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oquendo P., Alberta J., Wen D., Graycar J.L., Derynck R., Stiles "The platelet-derived growth factor-inducible KC gene encodes a secretory protein related to platelet alpha-granule proteins."; J. Biol. Chem. 264:4133-4137(1989).
PRINTS; PR00436; INTERLEUKIN8 PRINTS; PR00437; SMALLCYTKCXC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
MEDLINE=89139485; Pubmed=2917992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GRO_MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=96016008; PubMed=7561058;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE OF 1-10 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=129/SV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SCYB1 OR GRO1 OR GRO OR MGSA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cell Res.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              180:266-275(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          96
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                                                          QY
                                                                                                                                   DR FT FT SQ
                                                                                Matches
                                                                                         Query Match
Best Local 9
                                                                                                                                           DISULFID DISULFID
                                                                                                                                                                       PROSITE;
Cytokine;
SIGNAL
                                                                                                                                  SEQUENCE
                                                                                                                                                               CHAIN
                                                                                                                                                                                                       SMART; SM00199; SCY; 1.
 63
                    67
                                                           10 PVSMRLLAAALLLLL---LALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEK 66
                                        ω
                   MVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
  EVIATLKN----
                                       PATRSLLCAALLLLATSRLATGAPIANELRCQCLQTMAGIHLKNIQSLKVLPSGPHCTQT 62
                                                                                         Similarity
                                                                                                                                                                                            PS00471;
                                                                                                                                                                                  Growth factor;
                                                                                                                                  96 AA;
                                                                               Conservative
                                                                                                                                             25
33
GREACLDPEAPLVQKIVQ
                                                                                                                                                                                           SMALL_CYTOKINES_CXC;
                                                                                                                                  10254 MW;
                                                                                                                                             59
75
                                                                                         18.9%;
                                                                                                                                                                                 Inflammatory response; Signal
                                                                               18;
                                                                            Score 111; DB 1; L
Pred. No. 3.9e-05;
""matches 37;
                                                                                                                                           BY SIMILARITY.
BY SIMILARITY.
                                                                                                                                                               GROWTH REGULATED PROTEIN.
                                                                                                                                                                          PROBABLE
                                                                                                                                  4A52B5E5C38B45C2 CRC64;
                                                                                                  Length 96;
                                                                               Indels
                                                                              8
                                                                              Gaps
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2;

Search completed: January 29, 2002, 16:05:34 Job time: 120 sec

GenCore version 4.5 Copyright (c) 1993 - 2000 Comp

Compugen Ltd

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Result
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OM protein -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                               Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
506
332.5
1143.5
1128.5
128.5
97.5
93.5
87.6
86.5
88.5
88.5
88.5
76.5
76.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                 SPTREMBL_17:*

1: sp_archea:*
2: sp_bacteria
3: sp_fungi:*
4: sp_human:*
5: sp_inverteb:*
6: sp_mammal:*
7: sp_mhc:*
8: sp_organel1:
9: sp_phage:*
                                                                                                                                                                                                                        Query
 100.0

869.1

57.1

15.9

114.9

114.2

113.0

112.9

112.7
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11:
12:
13:
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Gapop 10.0 , Gapext 0.5
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587
1 MSLLPRRAPPVSMRLLAAAL.....TKRFIKWYNAWNEKRRVYEE 111
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                                                                                                                                                                                                                                                                                                           sp_rodent:*
sp_virus:*
sp_vertebrate:*
sp_unclassified:*
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sp_phage:*
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           Q9ERB1
O18209
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      O9btr1 homo sapien
O9ns21 homo sapien
O9ns21 homo sapien
O9jhh7 mus musculu
O9dfg4 brachydanio
O9dg18 gallus gall
O9ep62 rattus norv
O28724 oryctolagys
O98tq2 oncorhynchu
O9eqi5 mus musculu
O9eqi5 mus musculu
O9eqi6 mus musculu
O99me0 rattus norv
O73912 gallus gall
O9ptf8 brachydanio
O9xsx5 felis silve
O62764 equus cabal
O9erb1 mesocricetu
O18209 caenorhabdi
                                                                                                                                                                                                              Description
008889 cricetulus
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REGORDANCE OF THE PROPERTY OF

RESULT Q9NS21

O9NS21

Q9NS21

Q9NS21;
Q9NS21;
O1-OCT-2000 (TrEMBLrel. 15, Created)
O1-OCT-2000 (TrEMBLrel. 15, Last sequence update)
O1-UN-2001 (TrEMBLrel. 17, Last annotation update)
CHEMOKINE MIP-2 GAMMA.

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Query Ma Best Loo Matches 1/ 1 1 61 61 61	PRESULT 1 PART OF THE PART OF	02222222222222222222222222222222222222
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100.0%; Score 587; DB 4; Length 111; 2a1 Similarity 100.0%; Pred. No. 6.7e-58; 111; Conservative 0; Mismatches 0; Indels 0; Gaps 111; Conservative 0; Mismatches 0; Indels 0; Gaps MSLLPRRAPPVSMRLLAAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKY 60	PRELIMINARY; PRT; 11 (TrEMBLrel. 17, Created) 12 (TrEMBLrel. 17, Last sequ 13 (TrEMBLrel. 17, Last sequ 16 (TrEMBLrel. 17, Last sequ 16 (Human), 17 (Mast anno 17 (Mast) 18 (Human), 18 (Mast) 19 (74 12.6 394 5 044635 2.5 12.4 131 13 09PV19 2.5 12.4 356 11 088464 2.5 12.2 102 4 09C077 1.5 12.2 1417 5 09VSE1 71 12.1 1270 5 023658 0.5 12.9 101 13 093442 68 11.6 599 4 09P0W8 68 11.6 751 1 09WSY2 68 11.6 751 1 09WSY2 68 11.6 751 1 1 09WSY2 68 11.6 751 1 1 09WSY2 68 11.4 647 4 09WSY3 68 11.3 362 5 09WSY3 6.5 11.3 346 2 P70801 6.5 11.3 3228 5 093593 6.6 11.2 6815 5 0917U4 ALIGNMENTS ALIGNMENTS
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-OCT-2000 (TrEMBLrel. 17, Last annotation update)
01-JUN-2011 (TrEMBLrel. 17, Last annotation update)
B CELL AND MONONCYTE-ACTIVATING CHEMOKINE PRECURSOR
MNCB-6413, SIMILAR TO MUS MUSCULUS KIDNEY-EXPRESSED
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                    Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y., Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S., Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I., Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R., Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T., Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H., Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush Chemin L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio Yangha M., Wagner L., Washio
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Zhang W., Wan T., Yuan Z., Cao X.;
Zhang W., Wan T., Yuan Z., Cao X.;
"A novel chemokine gene, MIP-2 gamma.";
submitted (NOV-1998) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Osada N., Kusuda J., Hashimoto K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Watson J.D., Kumble K.D.;
"B cell and monocyte-activating chemokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN-BALB/CBYJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (KEC) MRNA) (KEC) (1200006123RIK PROTEIN).
SCYB14 OR BMAC OR 1200006123RIK.
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                                                                                                                                                                                                                                                        STRAIN=C57BL/6J; TISSUE=EMBRYONIC HEAD, AND LUNG; MEDLINE=21085660; PubMed=11217851;
                                                                                                                                                                                                                                                                                                                                                                          made by oligo-capping method.";
Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN-C57BL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (APR-1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=10090;
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                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                     isolation of full-length cDNA clones from mouse brain cDNA library
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Furuno
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Aono H., Baldarelli
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CHEMOKINE CXC
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Q9DFG4;
01-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Functional annotation of a full-length mouse cDNA collection.";
Nature 409:685-690(2001).
EMBL; AF144754; AAF66694.1; -.
EMBL; AB041614; BAA95097.1; -.
EMBL; AK014351; BAB29292.1; -.
EMBL; AK014351; BAB29292.1; -.
EMBL; AK004615; BAB29292.1; -.
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                                                                                                                                                                                                                                                            Brachydanio rerio (Zebrafish) (Zebra danio).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Ostariophysi;
Cypriniformes; Cyprinidae; Rasborinae; Danio.
                                                                                                                                                                                                                                                                                                                                               01-MAR-2001 (TrEMBLrel. 16, 01-MAR-2001 (TrEMBLrel. 16, 01-MAR-2001 (TrEMBLrel. 16,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                       during embryogenesis.";
Mech. Dev. 97:183-186(2000).
EMBL; AF279919; AAG09819.1;
                                                                                                                                                                distinctive expression patterns
                                                                                                                                                                                             Long
                                                                                                                                                                                                          SEQUENCE FROM N.A. PubMed=11025222;
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             14
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RLLAAALLLLLLALYTARVDGSKCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKMVIITTK 73
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MGI:1915056; 1200006123Rik.
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                                                       Similarity
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                                                                                                           12016 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23
99
                                                    57.1%;
54.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      86.2%;
94.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chemokine_IL8
                                        25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ω,.
                                                                                                                                                                                                                                                                                                                                                 Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                            Created)
                                                     Score 335; DB 13
Pred. No. 7.2e-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. 6.3
B; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 506;
Pred. No. 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     POTENTIAL.

B CELL AND MONONCYTE-ACTIVATING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CHEMOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97352E91FF7F46D6
                                                                                                             274BE255A74F98B5 CRC64;
                                        Mismatches
                                                                                                                                                                 novel CXC-type chemokine with the vestibulo-acoustic system
                                                                                                                                                                                                                                                                                                                                                                                                          100
                                                                 DB 13;
                                                                                                                                                                                                                                                                                                                                                                                                          ΑA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CRC64;
                                                                 Length 100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                          Indels
                                        0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                        Gaps
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Вb

RCSTAALFLLVIAVYSLNTEAYKCRCTRKGPKIRYIDVQKLEIKPKHPYCQEKMIFVTME 62

0,

Qγ

74 SVSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYE 110 :|||::|||::|| :|

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FTT DDR R TTTT
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                                                                                                                                                                                                                                                                                                                      RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAR-2001
01-MAR-2001
01-MAR-2001
          Signal.
                                                             Cytokine 10:169-174(1998).

EMBL; D87927; BAB12280.1; -.

EMBL; D87926; BAB1279.1; -.

InterPro; IPR001811; Chemokine_IL8.

InterPro; IPR001089; Small_cytokine_CXC.

Pfam; PF00048; IL8; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAR-2001 (Tremblrel. 01-MAR-2001 (Tremblrel. 01-JUN-2001 (Tremblrel. CINC-2 ALPHA PRECURSOR.
                             PROSITE; PS00471; SMALL_CYTOKINES_CXC; 1.
                                         PRINTS; PRO0437; SMAI
SMART; SM00199; SCY;
                                                                                                                                                          STRAIN-WISTAR; TISSUE-PERITONEAL CAVITY; MEDLINE-98236997; PubMed-9576061;
                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                   Q9EP62;
                                                                                                                         "Gene structure, cDNA cloning, and expression of rat cytokine-induced neutrophil chemoattractant 2 (GRO/CINC-2) gene.";
                                                                                                                                             Shibata F., Konishi K., Nakagawa H.;
                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                 NCBI_TaxID=10116;
                                                                                                                                                                                                                                 Rattus norvegicus (Rat).
                                                                                                                                                                                                                                                                                                Q9EP62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasiani
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gallus gallus (Chicken).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JUN-SUPPRESSED CHEMOKINE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q9DGL8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q9DGL8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9031;
                                                                                                                                                                                                                                                                                                                                                    61
                                                                                                                                                                                                                                                                                                                                                                    73 KSVSRYRG-QEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE 111
                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               63
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                                                                                                                                                                                                                                                                                                                                                                                          MKLLTAALLLLVIAMCLASAEGVKCKCSRKGPKIRFSNVRKLEIKPRYPFCVEEMIIVTL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                     60;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRELIMINARY;
   33
                                                     SMALLCYTKCXC.
\begin{array}{c} 32 \\ 101 \end{array}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11564 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                              54.9%;
60.0%;
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16,
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                                                                                                                                                                                                                                                     Last annotation update)
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Last annotation update)
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 POTENTIAL.
CINC-2 ALPHA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score
                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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No. 1.7e-28;
                                                                                                                                                                                                                                                                                               101
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RESULT
Q98TQ2
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Q28724
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Best Local S
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Best Local
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Q28724;
Q1-NOV-1996
Q1-AUG-1999
Q1-JUN-2001
Q98TQ2
                                                                                                                                                                                                                InterPro IPR001089; Small_cytokine_CXC.
pfam; pf00048; IL8; 1.
prints; pr000437; SMALLCYTKCXC.
SMARR; SM00199; SCY; 1.
proSITE; pS00471; SMALL_CYTOKINES_CXC; 1.
pS00471; SMALL_CYTOKINES_CXC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                   EMBL; U9$808; AAB93924.1; -.
EMBL; L28933; AAA66975.1; -.
HSSP; P19875; LQNK.
                                                                                                                                                                                                                                                                                                                                                  macrophages."
                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN-NEW ZEALAND WHITE;
Yoshimura T., Modi W.S.;
"Isolation of novel GRO genes, and a phylogenetic analysis
chemokine subfamily in mammals.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RPF2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GRO (PERMEABILITY FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                        Gene 151
                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=95129889; PubMed=7828903;
                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN-NEW ZEALAND WHITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted
                                                                                                                                                                                                                                                                                                                                                                           Martin T
                                                                                                                                                                                                                                                                                                                                                                                        Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9986;
                                                                                                                                                                                                                                                                                         [nterPro
                                                                                                                                                                                                                                                                                                                                                              'Cloning| of two rabbit GRO homologues and their expression in alveolar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                             69
                                                                        62 HCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61
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                                                                                                          AAP$GPRFLRTAMLLLLLLAASRRAAGAAALTELRCQCLQTVQGIHLKSIQSLKVLSPGP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PHCEEKMVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APP-TRRLLNAALLLLLLLMATSHQPSGTVVARELRCQCLKTLPRVDFENIQSLTVTPPG
                                                          HCAQTEVIATLKS-----GQEACLNPAAPMVKKFLQ
                                                                                                                               APPVSMRLLAAALLLLLLALYTARVDGS------KCKCSRKGPKIRYSDVKKLEMKPKYP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PHCTQTEVIATLKD-----GQEVCLNPQAPRLQKIIQ 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPVSMRLLAAALLLLLLALYTARVDGS-----KCKCSRKGPKIRYSDVKKLEMKPKY 60
                                                                                                                                                                                                                                                                                                                                                                                     ₹.C.,
                                                                                                                                                                     $imilarity
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                                                                                                                                                                                                                                                                                                                                       337-338(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                        OF 43-108 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                (MAR-1997) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                      IPR001811; Chemokine_IL8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6 (TrEMBLrel. 01, 09 (TrEMBLrel. 11, 17, 17)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         101 AA;
PRELIMINARY;
                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                        Goodman R.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11109 MW;
                                                                                                                                                                   21.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Last
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Created)
                                                                                                                                                       Score 128.5; DB 6
Pred. No. 7.9e-07;
5; Mismatches 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 143.5; DB 1
Pred. No. 1.6e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
                                                                                                                                                                                                                                                                                                                                                                                     Kajikawa O.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          D949D5712FE30909 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence update)
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97
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                                                            99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 11; Length 101;
                                                                                                                                                                                                                                                                                                                                                                                     Wong V.A., Mongovin S.M.,
                                                                                                                                                        38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31;
                                                                                                                                                        Indels
                                                                                                                                                                                Length 108;
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                                                                                                                                                        11;
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PRT;

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RESULT
Q9EQI5
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Q99J60
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Best Local
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Best Local
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Q9EQI5;
Q9EQI5;
01-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
NCBI_TaxID=8022;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Laing K.J., Zou J.J., Hirono I., Aoki T., Secombes .....
Tidentification and analysis of the interleukin 8 molecu
trout Oncorhynchus mykiss.";
Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases
EMBL; AJ279069; CAC33585.1;
EMBL; AJ279069; CAC33585.1;
SEQUENCE 97 AA; 10777 MW; 4EEB35A4EF9DACE2 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUN-2001 (TrEMBLrel.
01-JUN-2001 (TrEMBLrel.
01-JUN-2001 (TrEMBLrel.
PUTATIVE INTERLEUKIN 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; BC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q99J60;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUN-2001 (TremBLrel. 17, Created)
01-JUN-2001 (TremBLrel. 17, Last sequence update)
01-JUN-2001 (TremBLrel. 17, Last annotation update)
01-JUN-2001 (TremBLrel. 17, Last annotation update)
SMALL INDUCIBLE CYTOKINE B SUBFAMILY (CYS-X-CYS), MEMBER 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Strausberg R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=MAMMARY TUMOR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                        16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11 VSMRLLAAALLLLLLALYTARVDGS-----KCKC----SRKGPKIRYSDVKKLEMKP
                                                                                                                                                                                             61
                                                                                                                                                                                                                                                67
                                                                                                              10
                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                      1 MKSAVLFLLGIIFLEQCGVRGTLVIRNARCSCISTSRGTVHYKSLKDLKQFAPSPNCNKT 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9
                                                                                                                                                                                                                                                MVIITTKSVSRYRGQEHCLHPKLQSTKRFIK-WYNAWNEKRR 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MSIR-MSASLVVVLLALLTITEGMSLRGMGADLRCRCIETESRRIGKL----IKKVEMFP
                                                                                                                                                                                                                                                                                                                                                           LAAALLLLLLALY-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PSSHCRDTEIIATLSKS----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KYPHCEEKMVIIT-TKSVSRYRGQEHCLHPKLQSTKRFIK 97
                                                                                                                                                                                             EIIATLKN----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tted (FEB-2001) to the EMBL/GenBank/DDBJ databases BC003343; AAH03343.1; -.
NCE 126 AA; 14457 MW; 35892D1E125D1558 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 34.(
34; Conservative
(TrEMBLrel. 16,
                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                        PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRELIMINARY;
                                                                                                                                                                                             -GDQTCLDPDSANVKKLMKEWEKKINQKKK 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                        15.9%;
24.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16.6%;
34.0%;
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                                                                                                                                                                                                                                                                                                                                                        -TARVDGSKCKC-SRKGPKIRYSDVKKLEMKPKYPHCEEK
                                                                                                                                                                                                                                                                                                                                                                                                                    22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 93.5; DB 1
Pred. No. 0.0073;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 97.5;
Pred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                           113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 11;
                                                                                                                                                                                                                                                                                                                                                                                                                    40;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          126;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ij
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                                     RESULT
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                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                Zhang C., Poncz M., Thornton M.A., Kowalska A.;
Zhang C., Poncz M., Thornton M.A., Kowalska A.;
Zhang C., Poncz M., Thornton M.A., Kowalska A.;
"Localization of distal domains in the megakaryocyte-specific basic protein (pbp)/platelet factor 4 (PF4) gene locus.";
Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AF219112; AAG36786.1; -
EMBL; AF219112; AAG36786.1; -
EMBL; AF219173; BAB26811.1; -
EMBL; AF349465; AAK30162.1; -
EMBL; AF349465; AK430162.1; -
MGD; MGI:1922842; 2400003M24Rik.
InterPro; IPR001811; Chemokine_ILB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Gene Cloning and Functional Analysis of Novel Mouse Thymic Stromal Cells.":
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
THYMUS CHEMOKINE 1 (2400003M24RIK PROTEIN) (PLATELET BASIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=C57BL/6J; TISSUE=EMBRYONIC MEDLINE=21085660; PubMed=11217851;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=THYMUS;
Gong S.-Y., Ch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2400003M24RIK.
                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                              PRINTS; PR00437; SMALLCYTKCXC SMART; SM00199; SCY; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hayashizaki Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kawai J., Shinagawa A., Shibata K., Yoshino M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=10090
                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nature 409:685-690(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Functional annotation of a full-length mouse
                                                                                                    83
                                                                                                                                                                                                                      15
                                                                                                                                     68 VIITTKSVSRYRGQEHCLHPKLQSTKRFI 96
                                                                                                                                                                               23
                                         11
                                                                                                                                                                                                                  LLAAALLLLLLALYTA-RVDGS-----KCKCSRKGPKIRYSDVKKLEMKPKYPHCEEKM 67
                                                                                                                                                                               LLLLGLILVALAPLTAGKSDGMDPYIELRCRCTNTISGIPFNSISLVNVYRPGVHCADVE 82
                                                                                                    VIATLKN----
                                                                                                                                                                                                                                                                                                                                                                                                                       PF00048;
                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                          IPR001089; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (DEC-1999)
                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                   IL8;
                                                                                                                                                                                                                                                                                                                                                              AA:
                                                                                                  -GQKTCLDPNAPGVKRIV 106
                                                                                                                                                                                                                                                                                                                                                              12251 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to the
                                                                                                                                                                                                                                                                               15.2%;
29.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Small_cytokine_CXC
                                                                                                                                                                                                                                                             15;
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                                                                                                                                                                                                                                                                               Score 89;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                              66F93D3F7D7862C8 CRC64;
                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STEM CELLS
                                                                                                                                                                                                                                                                             DB 11; Length 113; 0 021;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Muridae;
                                                                                                                                                                                                                                                               36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cDNA collection.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Guo J.-Z.;
Chemokines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Itoh M.,
                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ishii Y.,
Fukuda S.,
mmanaka I.,
Saito R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Secreted
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01-AUG-1998
01-JUN-2001
                    HSSP; P02775; lTVX.
InterPro; IPR001811; Chemokine_IL8.
InterPro; IPR001089; Small_cytokine_CXC.
Pfam; PF00048; IL8; 1.
PRINTS; PR00437; SMALLCYTKCXC.
SMART; SM00199; SCY; 1.
                                                                                                           Hughes S.M., Bumstead N.;
"Mapping of a second ELR CXC chemokine to chicken chromosome four.";
Submitted (JUN-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; Y14971; CAA75212.1; -.
                                                                                                                                                                                                  MEDLINE=20170941; PubMed=10704244; Sick C., Schneider K., Staeheli P., Weining K.C. "Novel chicken CXC and CC chemokines.";
         PROSITE;
                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                             Cytokine 12:181-186(2000).
                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
TISSUE=MACROPHAGE LIKE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Cloning and Characterization of a Novel Gene Coding the Gene to MTCK1.";
Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                        Sick C
                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                             Gallus gallus (Chicken).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             073912;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           073912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUN-2001 (TrEMBLrel. 17, 01-JUN-2001 (TrEMBLrel. 17, 01-JUN-2001 (TrEMBLrel. 17,
                                                                                                                                                                                                                                                                        Submitted (AUG-1997) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9031;
                                                                                                                                                                                                                                                                                                                                                                   Archosauria; Aves; Neognathae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RTCK1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CXC CHEMOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    66
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                                                                                                                                                                                                                                                                                                                                                                                                                      AUG-1998 (TremBLrel. 07, Created)
AUG-1998 (TremBLrel. 07, Last sequence update)
JUN-2001 (TremBLrel. 17, Last annotation updat)
PROTEIN PRECURSOR (CXC CHEMOKINE K60)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KMVIITTKSVSRYRGQEHCLHPKLQSTKRFIK 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SMRLLAAALLLLLLALYTARVDGS-----KCKCSRKGPKIRYSDVKKLEMKPKYPHCEE 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VEVIATLKN-----GKEVCLDPTAPMIKKIVK 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NLQVLLSLLLVALVPLAVGKPDNNEPFIELRCRCTNTLSGIPLNSISRVNVFRPGAHCDN 78
                                                                                                  AF277660; AAF86485.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tted (FEB-2001) to the EMBL/GenBank/DDBJ databases AF349115; AAK30166.1; -. NCE 111 AA; 12159 MW; 8230226BC7C60A55 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
22; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gong S.,
      PS00471; SMALL_CYTOKINES_CXC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14.9%;
23.9%;
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        No. 0.03;
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Q9XSX5
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Best Local
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01-NOV-1999 (TrEMBLrel.
01-NOV-1999 (TrEMBLrel.
01-JUN-2001 (TrEMBLrel.
SEQUENCE FROM N.A.
Straubinger A.F., Simpson K.W.,
"Feline interleukin-8 mRNA.";
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CHAIN
SEQUENCE
                                                      Eukaryota;
Mammalia;
                                                               Felis silvestris catus (Cat).
Eukaryota; Metazoa; Chordata;
                                                                                                                                             Q9XSX5
                                                                                                                                                                                                                                                                                                                                                          "Chemokines and chemokine receptors in zebrafish."; Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases. EMBL; AF$01450; AAF17560.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Brachydanio rerio (Zebrafish) (Zebra danio).
Eukaryda; Metazoa; Chordata; Cranilata; Verte
Actinopterygii, Neopterygii; Teleostei; Eutel
Cypriniformes; Cyprinidae; Rasborinae; Danio.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAY-2000 (TrembLrel. 13, 01-MAY-2000 (TrembLrel. 13, 01-JUN-2001 (TrembLrel. 17, CC CHEMOKINE CCL1.
                                        NCBI_TaxID=9685;
                                                                                       INTERLEUKIN-8
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Rollins B.J.;
                                                     Eutheria; Carnivora;
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                                                    Craniata; Vertebrata;
; Fissipedia; Felidae;
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83725AB91FE18E3D CRC64;
          Straubinger R.K.;
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Best Local Similarity
Matches 30; Conserv
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Best Local Similarity 28.0
Matches 28; Conservative
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Q68398;
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EMBL; AF158598; AAD40323.1; -.
HSSP; P10145; 1IKM.
InterPro; IPR001811; Chemokine_ILB.
InterPro; IPR001089; Small_cytokine_CXC.
Pfam; PF00048; ILB; 1.
PF1MTS; PR00048; ILB; 1.
PRINTS; PR000437; SMALLCYTKCXC.
SMART; SM00199; SCY; 1.
PROSITE; PS00471; SMALL_CYTOKINES_CXC; 1.
PROSITE; PS00471; SMALL_CYTOKINES_CXC; 1.
                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.

STRAIN-TOLEDO;

MEDLINE-96099416; PubMed-8523595;

Cha T.A., Tom E., Kemble G.W., Duke G.M., Mocarski E.S., Spaete R.R.;

Cha T.A., Tom E., Kemble G.W., Duke G.M., Mocarski E.S., Spaete R.R.;

"Human cytomegalovirus clinical isolates carry at least 19 genes not

"found in laboratory strains.";

J. Virol. 70:78-83(1996).

EMBL; U33331; AAA85885.1; -.

EMBL; U33331; AAA85885.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human cytomegalovirus.
Viruses; dsDNA viruses, no RNA stage; Herpesviridae; Betaherpesvirinae; Cytomegalovirus.
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
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                                                                                                                                            83
                                                                                                                                                                                                         13 MRLLAAALLLLLALYTARVDGS--KCKC-SRKGP--KIRYSDVKKLEMKPKYPHCEEKM 67
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                                                                                                                                          VIITTKSVSRYRGQEHCLHPK-----LQSTKRFIKWYNAWNEKR 106
                                                                                                                                                                                  MRLIFGALIIFLAYVYHYEVNGTELRCRCLHRKWPPNKIILGNY-WLHRDPRGPGCDK---
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27.5%;
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%; Pred. No. 0.088;
15; Mismatches
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P. 15

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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  January 30, 2002, 02:06:02; Search time 2414.34 Seconds (without alignments) 11513.601 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

RESULT 1 AXI 36401 LOCUS DEFINITION SE ACCESSION AX VERSION AX VERSION AX KEYWORDS hu ORGANISM HO EAU AUTHORS HA TITLE SE JOURNAL HA FEATURES SOURCE HE FEATURES SOURCE HE		44 44 44 44 44 44 44 44 44 44 44 44 44	20 435.6 21 386 22 368 22 367.2 24 355.4 25 354.6 27 299.8 28 150.8 29 150.8 20 122.8 20 31 76.4 21 76.4	9 976. 0 976. 1 872. 2 872. 2 873. 3 733. 4 727. 5 709. 5 709. 5 58 8 550.	164 143 117 117 117 117 117 1109	Result No. Score
man. mo sapiens karyota http://www.new.ar. man. mo sapiens protent hetheria; Primates; a,T., Isogai,T., Nishikawa, a,T., Isogai,T., Nishikawa, protein or membrane pretent: EP 1067182-A 323 10-J, lix Research Institute (JP) Location/Qualifiers 11677 /organism="Homo sapi.	ALIC	N N N O N N O N O O N N O	25.9 461 6 AR071 22.9 386 9 AF073 21.8 752 10 AF25 21.8 533 10 AF35 21.1 504 10 AF15 21.0 354 9 HUMADD 21.0 354 9 HUMADD 21.0 354 9 FUMADD 21.0 354 PUMADD 21.0 354	57.9 198 57.9 199 51.8 199 51.8 199 43.5 186 43.2 166 42.1 188 34.5 58 34.5 58	97.4 1677 6 85.1 1503 9 70.0 108123 9 69.9 134506 9 69.8 79970 2 69.8 159420 2 69.1 81811 2 64.8 1630 6	Query Match Length DB ID
PAT 1067182. PAT ; Craniata; Vertebrata ; Catarrhini; Hominida ; Catarrhini; Sugiyam ne protein ne protein JAN-2001; s	ALIGNMENTS	AC0910; AC0910; AF3108 BC00066; AC0253 AC11106; AC0254 AC00344 BC0025; AC03622 AC01634 AC08711;	AR0712; AF0732; AF0732; AF952; AF9863; AF9863; AF9863; AF2858;			Descri
30-MAY-2001 ; Euteleostomi; e; Homo.		sequence 14 23 Homo sapi 84 Plasmodiu 96 Dictyoste 08 Homo sapi 18 Homo sapi 18 Homo sapi 19 Homo sapi 10 Homo sapi	equence omo sap Mus mus Mus mus omo sap equence equence anlus mus allus g anio re GC-1304 AF-209	equencequencequencequencequencequencemus mu Mus mu Mus mu Mus mu omo sa equencequencequenc	01 Sequence 03 Homo sapi 06 Homo sapi 38 Homo sapi 28 Homo sapi 17 Homo sapi 181 Homo sapi 19 Sequence	cription

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ACCESSION REFERENCE SOURCE KEYWORDS VERSION ORGANISM

Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. AF144103.1 Homo sapiens AF144103 AF144103 GI:5059165 NJAC 1503 bp mRNA protein (NJ (NJAC) mRNA, complete cds

JOURNAL expression Unpublished Identification of a novel chemokine family member with altered expression in human head and neck squamous cell carcinoma

TITLE

(bases 1 to 1503)

gene
/map="5q25-q31" /tissue_type="normal oral epithelial" 11503 /gene="NJAC"

KEYWORDS SOURCE ORGANISM

Homo sapiens chromosome AC034206 AC034206.4 GI:14518406

108123 bp

5 c clone

PRI CTC-321K16,

20-JUL-2001 complete sequence

REFERENCE

Human. Hagiens
Human sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
1 (bases 1 to 108123)

Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.

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Direct Submission

Submitted (05-APR-2000) Production Sequencing Facility,

Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA
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 {\tt Direct} Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Quality: Phrap Quality >=40 99.5% of Sequence; Estimated Total Number of Errors is 0.2. STS Content: SHGC-130469 G59717.
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On Jun 21, 2001 this sequence version replaced gi:9256720.
Draft Sequence Produced by DOE Joint Genome Institute
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Homo sapiens chromosome
Unpublished
              Comparison Analysis (SCAN)
                            Ricke, D.O.
Large Scale Sequence Analysis and
                                                                      Unpublished
                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria, Primates; Catarrhini; Hominidae; Homo.
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BAC clone 7g12 (LBNL H126), complete
                 System
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2889360995 /organism="Homo sapiens"	2450724641 /rpt_family="MER7"	complement(2337423671) /rpt family="A u"	complement(2362822797) /rnt family="MTP"		20251. 2019	/rpc_ramily="Alu" complement(19591 .19738) /rnf family="MTP"	/rpc_ramily="L1" 1516315487	1477914955	/rpt_tamily="MER42" 1217312472	/rpt_family="Alu" 1067410876	/rpt_tamily="MER42" 1034610624	7 PC_IdIII1Y="MER42" 974510340	/IDC_IdIIIII/V="MIX"	complement(5036, 5148)	complement(39404205)	8832994)	2 excellent exco frame	ement(="GRAI	/cnromosome="5" /map="5p"	sm="Homo sapiens" f="taxon:9606"	/chromosome="5" join(129755,7816982395,123646134506)	:9606"	Location/Qualifiers 1. 134506	it Genome Institute.	HRL-1, LS-3, MS M888, LOS Alamos, NN	bmission (20-OCT-1998)	es 1 to 134506)	Direct Submission Submitted (13-OCT-1998) Los Alamos National Laboratory, DOE Joint Genome Institute, TA43, HRL-1, LS-3, MS M888, Los Alamos, NM 87545,	CA 94720, U.S.A. 1 to 134506)	Institute, Lawrence Berkeley National Laboratory, MS 74-157,	3 (bases 1 to 134506) Kimmerly, W., Bondoc, M., Cheng, J., Connolly, K.S., Gunning, K.M., Kadmer, K., Miguel, T., Miller, C., Pitluck, S., Pollard, M., Rojeski, H., Subramanian, S. and Martin, C.H.	

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complement(77047...77436)
/rpt_family="MLT1"
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/note="GRAIL 2 excellent exon,
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complement(82876. .82937)
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complement(72274.
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                                                                Submitted (06-OCT-1999) Production Sequencing Facility, DOE Joint Genome Institute, .2800 Mitchell Drive, Walnut Creek, CA 94598, USA On Nov 15, 2000 this sequence version replaced gi:6604400.
                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 79970)
DOE Joint Genome Institute.
Sequencing of Human Chromosome 5
                                                                                                                                                                                                                                                                                                                                                AC011428 79970 bp DNA
Homo sapiens chromosome 5 cl
SEQUENCE, 4 unordered pieces
             Center: Joint Genome Institute
Center Code: JGI
                                                                                                                2 (bases 1 to 79970)
DOE Joint Genome Institute.
Direct Submission
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AC011428.3 GI:11178060
                                                                                                                                                               Unpublished
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Web site:
                                                  ---Genome Center
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http://www
                                                                                                                                                                                                                                                                                                     HTGS_DRAFT
jgi.doe.gov
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Consensus quality: 78432 bases at least 020
Consensus quality: 78894 bases at least 020
Consensus quality: 78894 bases at least 020
Estimated insert size: 72000; pulse field gel estimation
Estimated insert size: 79670; sum-of-contigs estimation
Ouality coverage: 7.19 in 020 bases; pulse field gel estimation
Ouality coverage: 6.49 in 020 bases; pulse field gel estimation.
* NOTE: This is a 'working draft' sequence. It currently
* consists of 4 contigs. The true order of the pieces
is not known and their order in this sequence record is
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18934 c 18691 g 21162
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3324: gap of unknown length
13887: contig of 10563 bp in length
13987: gap of unknown length
32432: contig of 18445 bp in length
32532: gap of unknown length
79970: contig of 47438 bp in length.
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 Center: Join
Center Code:
                                           Direct Submission
Submitted (03-AUG-1999) Production Sequencing Facility, DOE Joint
Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USI
On Apr 20, 2001 this sequence version replaced gi:11178048.
                                                                                                                                                       1 (bases 1 to 159420)
DOE Joint Genome Institute.
Sequencing of Human Chromosome
                                                                                                                                                                                        Eukharyota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 159420)
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DOE Joint Genome Institute.
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HTG; HTGS_PHASE1; HTGS_DRAFT;
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Consensus quality: 142845 bases at least Q40 consensus quality: 150910 bases at least Q30 consensus quality: 152756 bases at least Q30 consensus quality: 152756 bases at least Q30 estimated insert size: 80000; pulse field gel estimation Estimated insert size: 156920; sum-of-contigs estimation Quality coverage: 8.41 in Q20 bases; pulse field gel estimation Quality coverage: 4.29 in Q20 bases; sum-of-contigs estimation.*
*NOTE: This is a 'working draft' sequence. It currently consists of 26 contigs. The true order of the pieces time that the property of the pieces to the pieces the pieces to the pieces the pieces the pieces to the pieces the pieces to the pieces to the pieces the pieces to the pieces the pieces to the piec
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                                                                                                                   Consensus quality: 74154 bases at least Q40
Consensus quality: 79224 bases at least Q20
Consensus quality: 80246 bases at least Q20
Estimated insert size: 76000; pulse field gel estimation
Estimated insert size: 176000; pulse field gel estimation
Quality coverage: 5.65 in Q20 bases; pulse field gel estimation
Quality coverage: 5.28 in Q20 bases; sum-of-contigs estimation.

* NOTE: This is a 'working draft' sequence. It currently

* consists of 11 contigs. Gaps between the contigs

* are represented as runs of N. The order of the pieces

* is believed to be correct as given, however the sizes

* to the gaps between them are based on estimates that have
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (22-APR-2000) Production Sequencing Facility, DOE Joint Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA on Sep 2, 2000 this sequence version replaced gi:7636394.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Center: Joint Genome Institute Center Code: JGI
                                                                                                                                                                                                                                                                                                                                              Summary Statistics
                                                                                                                                                                                                                                                                                                                                                                               Center clone name: XXp1-766F5
                                                                                                                                                                                                                                                                                                                                                                                            Project Information
Center Project Name: 1189125,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Web site: http://www.jgi.doe.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens chromosome 
11 ordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOE Joint Genome Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                by the finished sequence as soon as it is available the accession number will be preserved.

1 3978: conting of 3978 bp in length
3979 4078: gap of unknown length
                                                                                       This sequence will be replaced
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Matches 1189;
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                                                                                                                                                                                                                                                                                                                                                                                     1082 tttgpttgccagaaaaatgtgcttcattcccccctggttaatttttacacaccctaggaa 1141
                                            1022 cccckgtagccgggcaagcaggagcaggtctctctgcatctgttctctgaggaactcaag 1081
                                                                                                                                           842 acatctgagctttataagcgcccgggaggaacaatgagcttggtggacacatttcattgc
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                             CCCCFGTAGCCGGGCAAGCAGGAGCAGGTCTCTCTGCATCTGTTCTCTGAGGAACTCAAG
                                                                                      GCTGCCACGGGCTCTCCTGGGCCTTATGGCCGGTCACAGCCTCAGTGTGACTCCACAGTGG
                                                                                                    gctgbcacgggctctcctgggcttatggccggtcacagcctcagtgtgactccacagtgg
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                                                                                                                                                                                                                                                                  GAAAACTCATGCCTTTCCTTTTTAAAAAATGCTTTTTTGTATTTGTCCATACGTCACTAT
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/db_xref="taxon:9606"
/chromosome="5"
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99.4%;
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81811: contig of 920 bp in length
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JOURNAL
                                   Matches 1094;
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            aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaagcctttcttctcacaggcat
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Sequence 59 :
AX013119
AX013119.1 (
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                               Human nucleic acid sequences of endometrium tumour tissue Patent: WO 9954461-A 59 28-OCT-1999; SCHMITT ARMIN (DE); SPECHT THOMAS (DE); DAHL EDGAR (DE); BERND (DE); ROSENTHAL ANDRE (DE); METAGEN GES FUER GENOMF (DE); PILARSKY CHRISTIAN (DE)
                                                                                                                                                                                                                             Schmitt,A., Specht,T., Dahl,E., Hinzmann,B.,
Pilarsky,C.
                                                                                                                                                                                                                                                  (bases 1 to 1630)
                                   Conservative
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                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
1 392 c 327 g 43
                                                                                                                                                      Location/Qualifiers
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99.7%;
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                                   0;
                                             Score 1092.2; DB 6;
Pred. No. 8.3e-201;
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                                                                     ctgcatctgttctctgaggaactcaagtttggttgccagaaaaatgtgcttcattccccc 1114
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Patent: WO 9955888-A 77 04-WOV-1999;
SCHMITT ARMIN (DE); SPECHT THOMAS (DE); DAHL E
BERND (DE); ROSENTHAL ANDRE (DE); METAGEN GES
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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Sequence 77
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Human nucleic acid sequences obtained from pancreas Patent: WO 9955858-A 77 04-NOV-1999; SCHMITT ARMIN (DE); SPECHT THOMAS (DE); DAHL EDGAR (
                                      Pilarsky,C
                                            1 (bases 1 to 1962)
Schmitt,A., Specht,T., Dahl,E., Hinzmann,B.,
                                                                         Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
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                                              CCCAACCTGAGGATTTCTGAAAGGTTCACAGGTTCAATATTTAATGCTTCAGAAGCATGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
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97.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 872.6; DB 6;
Pred. No. 1.9e-158;
0; Mismatches 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             544 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PAT
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JOURNAL REFERENCE
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AB041614
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                                                                                                                                                                                                                                                                                    Direct Submission
Submitted (12 APR-2000) to the DDBJ/EMBL/GenBank databases.
Submitted (12 APR-2000) to the DDBJ/EMBL/GenBank databases.
Katsuyuki Hashimoto, National Institute of Infectious Disea
Division of Genetic Resources; 23-1, Toyama 1-chome, Shinju
Tokyo 162-8640, Japan (E-mail:khashi@nih.go.jp,
URL:http://www.nih.go.jp/yoken/genebank/,
Tel:81-3-5285-1111(ex.2120), Fax:81-3-5285-1181)
URL: http://www.nih.go.jp/yoken/genebank/
Tel:81-3-5285-1111(ex.2120), Fax:81-3-5285-1181)
URL: http://www.nih.go.jp/yoken/genebank/
                                1st strand cDNA was primed with an oligo(dT) primer [ATGTGGCCTTTTTTTTTTTTTTTTTTT]; double-stranded cDNA was synthesized using specific 5' and 3' primers and amplified by PCR. The PCR product was digested with SfiI and size selection was performed to exclude fragments <1.5kb.The SfiI-digested PCR product was cloned into distinct DraIII sites of pME188-FL3. XhoI sites just outside the DraIII sites can be used to isolate the cDNA insert. Library was constructed by Sugano et al.(University of Tokyo, Institute of Medical Science). Custom primer used for sequencing (5' end primer ICTTCTGCTCTTAAAACCTGCG); 3' end primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2000)
2 (bases 1 to 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Osada, N.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  made by oligo-capping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       isolation of full-length cDNA clones from mouse brain cDNA library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hashimoto, K
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                     CGACCTGCAGCTCGAGCACA] ).
  part of this sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        s (full insert sequence).
s musculus (strain:C57BL) adult female cDNA to mRNA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      method
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is reported in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            clone MNCb-6413, similar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mncb clone: MNCb-6413.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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AU035952
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to Mus musculus
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                                                                                                                                        accaagcgcttcatcaagtggtacaacgcctggaacgagaagggcagggtctacgaagaa
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                                                                                    ctttctcacaggcataagacacaaattatatattgttatgaagcacttttttaccaacggt
                                                                                                                                                                                                                         ACCAAACGCTTCATCAAGTGGTACAATGCCTGGAACGAGAAGCGCAGGGTCTACGAAGAA
                                                                                                                                                                                                                                                                        GTGGACGGGTCCAAGTGTAAGTGTTCCCCGGAAGGGGCCCCAAGATCCGCTACAGCGACGTG
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gtgctccagacttcatcacaggctgcttttt-----
                                                                                                                          TTTGCAGATT - -
                                                                                                                                                                                      tagggtgaaaaacctcagaagggaaaactccaaaccagttgggagacttgtgcaaaggac
                                                                       TAGGGTGGACGATCATGGAAAAGAAAACTCCAGGCCAGTTGAGAGACTTCAGCAGAGGAC
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KPKYPHCEEKMVIVTTKSMSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRVYEE"
a 520 c 499 g 382 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /strain="C57BL"
/db_xref="taxon:10090"
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/db_xref="GI:7670492"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /codon_start=1
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/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="MNCb-6413"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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Pred. No. 1.3e-131;
0; Mismatches 361;
                                                                                                                          -----AAAATAAAAGCCCTTT
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atcaaaaaggggaaaa
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RESULT 3 AF144754

DEFINITION

AF144754 Mus musculus (Bmac) mRNA,

1663 bp mI B cell and mc complete cds.

mRNA

and mononcyte-activating chemokine

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/translation="MRLLAAALLLLLLALCASRVDGSKCKCSRKGPKIRYSDVKKLEM
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Query Match Best Local

Local Similarity

42.18; 70.98;

Score 709.6; DB 10; Pred. No. 5.2e-127;

Length 1884;

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                                                                                                                                                                                                                                                                                                             Submitted (06-OCT-1999) Genetics, Road, Piscataway, NJ 08854-8082, Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                   Wang,L., Deng,L., Raikwar,N., Sahota,A. and Tischfield,J.A. Identification of a kidney-expressed chemokine (KEC), a member of the CXC family, that is selectively elevated in aprt knockout mice
                                                                                                                                                                                                                                                                                                                                                                    Wang, L., Deng, L., Direct Submission
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          /product="kidney-expressed chemokine CXC"
/protein_id="AAF03753.1"
/db_xref="GI:6103733"
/translation="MRILAAALILLLIALCVRALDGSKCKCSRKGPKIRYSDVKKLEM
KPKYPHCEEKMVIVTTKSMSRYRGQEHCLHPKLQSTKRFIKWYNAWNEKRRYYEE"
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/strain="C3H"
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cacagctgccacgggctctcctgggcttatggccggtcacagcctcagtgtgactccaca 1017
                                                CAGCACTACTGCTCCGCTCCAGGCTTACAAAGCTTCCGCTCAGAGAGCCTGGCGGCTCTG
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gaaagtaataaaatatatttgaaatgtaaaaaaaaa	tatgtacatgtcagaaaccattagcattgcatgcaggtttcatattctttct	tocacacactgcccccagtatatgccgcattgtactgctgtgttatatgc 1585	gctaagatgaccatgcgccctttcctctgtacatatacccttaagaacgccccc- 1534 	atatatttgtgattccccatgtaattcttcaatgttaaacagtgcagtcctctttcgaaa 1480	aaccttcaaagcatgtttctttccctcaccacaacagaacatgcagtactaaagca 1420	atatatgaatacatgcgcaatacacagctacagacacacattctgttgacaagggaa 1364 	Catgtgaggttcccaacactgtcagcaaaaccttaggagaaaacttaaaa 1307	ctt-tcccaacctgaggatttctgaaaggttcacaggttcaatatttaatgcttcagaag 1256 	99aaacatttccaagatcctgtgatggcgagacaaatgatccttaaagaaggtgtgggggt 1197 	caagititggtigccagaaaaatgigcitcaticccccctggitaatitittacacacccta 1137	9t99cccct9ta9cc999caa9ca99a9ca99tctctctgcatct9ttctctgaggaact 1077

Search completed: January 30, 2002, 04:09:05 Job time: 7383 sec

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Oligonucleotide D1	AAF58257	22	936		89.8	45
	AAF58254	22	936		89.8	44
Oligonucleotide D1	AAF58252	22	936		89.8	43
#19472 used	AAI50786	22	136		114.8	42
#6299 used	AAI37613	22	474		115.8	41
e microglia a	AAS06432	22	201	8.6	144.4	40
O	AAC99731	22	234		214.8	39
cDNA fragme	AAZ61798	21	234		214.8	38
cell c	AAC99732	22	234		234	37
CDNA	AAZ61799	21	234		234	36
ene signa	AAT25671	16	276		241.3	35
ed partial	AAV43787	19	276		248.8	34
ecreted p	AAC08310	21	275		264	33
l al	AAV43786	19	374		267.8	32
n ge	AAT21780	16	359		277.8	31
al rodent ch	AAV43785	19	375		279.8	30
	AAC99730	22	288	17.	288	29
Human huKS1 cDNA.	AAD10141	22	288	17.	288	28
	AAZ61797	21	288	17.	288	27
	AAZ08969	20	478	19.	322.2	26
	AAX89708	20	336	19.	336	25
endometrium	AAZ42039	20	1630	20.	352.8	24
	AAC99576	22	766	25.	426	23
Ë	AAZ61643	21	766	25.	426	22
n chemokine a	AAT86316	18	461	25.	435.6	21
Skin cell cDNA, SE	AAC99723	22	764	26.	448	20
Mouse muKS1 cDNA.	AAD10140	22	764	26.	448	19
⊂ −	AA261790	21	764	26.	448	18
specific f	9420	22	584		550.4	17
Human colon cancer	9819	21	726		10	16
er specific f	AAF94062	22	599		580	15
okine-1 par	AAZ08964	20	1372		60	14
ne neokine-1	9	20	1656	42.8	720.6	13
Skin cell cDNA, SE	AAC99808	22	99		727.2	12

ALIGNMENTS

Human PRQ273 nucleotide sequence. standard; cDNA; 1685 (first entry) ВP

Human; PRO; EST; expressed sequence tag; PCR primer; hybridisation; probe; blood coagulation disorder; cancer; cellular adhesion disorder; secreted protein; transmembrane protein; ss.

99WO-US05028

98US-0077632. 98US-0077641.

98US-0077649. 98US-0077791. 98US-00778004. 98US-0040220. 98US-0078886. 98US-0078936. 98US-0078936. 98US-0078939.

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PR 25-MAR-1998; 98US-0079656.
PR 27-MAR-1998; 98US-0079664.
PR 27-MAR-1998; 98US-0079664.
PR 27-MAR-1998; 98US-0079786.
PR 27-MAR-1998; 98US-0079728.
PR 30-MAR-1998; 98US-0079728.
PR 31-MAR-1998; 98US-0080105.
PR 30-MAR-1998; 98US-0080107.
PR 31-MAR-1998; 98US-0080107.
PR 31-MAR-1998; 98US-0080127.
PR 01-APR-1998; 98US-0080327.
PR 01-APR-1998; 98US-0080327.
PR 01-APR-1998; 98US-0080327.
PR 01-APR-1998; 98US-0081070.
PR 08-APR-1998; 98US-0081175.
PR 15-APR-1998; 98US-0081170.
PR 08-APR-1998; 98US-0081170.
PR 15-APR-1998; 98US-00811203.
PR 15-APR-1998; 98US-0081207.
PR 15-APR-1998; 98US-0081207.
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PR 29-APR-1998; 98US-0083336.
PR 29-APR-1998; 98US-0084663.
PR 29-APR-1998; 98US-0085573.
PR 29-APR-1998; 98US-0085593.
PR 29-APR-1998; 98US-0085593.
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30-JUL-1998;
11-SEP-1998;
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DB; AAY41739.
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                                          gcgcagggtctacgaagaatagggtgaaaaacctcagaagggaaaactccaaaccagttg
                                                                                     gcaccccaagctgcagagcaccaagcgcttcatcaagtggtacaacgcctggaacgagaa
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                            gcgcagggtctacgaagaatagggtgaaaaacctcagaagggaaaactccaaaccagttg
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30-NOV-1999

02-DEC-1999

02-DEC-1999

16-DEC-1999

16-DEC-1999

30-DEC-1999

equence tag) sequences which encode secreted or transmembrane PRO polypeptides. The PRO polynucleotides and polypeptides have cytostatic activity. The polynucleotides and polypeptides can be used for detecting the presence of PRO polypeptides in samples, for linking bioactive molecules to cells and for modulating biological activities of cells, using the polypeptides for specific targeting. The polypeptide targeting can be used to kill the target cells, e.g. for the treatment of cancers. The polypeptide pairs provide specific targeting of bioactive molecules
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Shelton
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Goddard
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                                                                                                                                            Novel PR\Diamond polypeptides and polynucleotides used in de to target bioactive molecules to specific cells, and cellular activities -
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Filvaroff E,
Godowski PJ,
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2000WO-US00277.
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Stewart
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Grimaldi CJ, Gurne,
pier MA, Pan J, Pao
Tumas D, Williams |
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Gurney AL, Hillan J
J, Paoni NF, Roy MA
Llliams PM, Wood WI;
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standard; cDNA;

(first entry)

Human PRO273 protein UNQ240 encoding cDNA SEQ ID

NO:45

ARESULT AACSSE ID AACSS XX AACSS XC AACSS XX AACSS XX Huma DE Huma XX Huma CX dermatological; antiarthritic; antirheumatic; immunosuppressive; haemostatic; antithyroid; antidiabetic; nootropic; neuroprotective; antianaemic; hepatotropic; virucide; antipsoriatic; antiallergic; antiasthmatic; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; spondyloarthropathy; systemic sclerosis; sarcoidosiidiopathic inflammatory myopathy; Sjogren's syndrome; thyroiditis; systemic vasculitis; autoimmune haemolytic anaemia; diabetes mellit immune related disease; diagnosis; antiinflammatory; thrombocytopaenia; immune-mediated sarcoidosis; cardiant;

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anaemia,

selected from systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis, spondyloarthropathies, systemic sclerosis, idiopathic inflammatory myopathies, Sjogren's systemic systemic vasculitis, sarcoidosis, autoimmune haemolytic,

immune-mediated renal disease, demyelinating diseases of the central and peripheral nervous systems, hepatobiliary diseases, inflammatory

other;

used

Indels Length

0;

Gaps

0;

autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus,

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20-APR 1999
28-APR 1999
04-MAY 1999
02-JUN 1999
23-JUN 1999
26-JUL-1999
26-JUL-1999
27-JUL-1999
28-JUL-1999
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29-OCT 1999
15-SEP 1999
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31-FEB 20000
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Kabakoff RC,
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2000WO-US00277.
2000WO-US00376.
2000WO-US03565.
2000WO-US04341.
2000WO-US04342.
2000WO-US04414.
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99WO-US28313.
99WO-US28409.
99WO-US28301.
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99WO-US28554.
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Lu Y, Pan
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99US-01456918
99US-0146222
99WO-US201111
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Sixty four PRO polypeptides, useful in the diagnosis and treatment immune related disorders, e.g. systemic lupus erythematosis, rheuma arthritis, osteoarthritis, thyroiditis and diabetes mellitus -
             The present invention describes sixty four human PRO be used in the treatment of immune related diseases. proteins, anti-PRO antibodies, agonists and antagonists.
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cgcagccggagccagcagagccggaaggcgcgccccgggcagagaaagccgagcagagct

Matches 1660; Query Match Best Local Similarity

Conservative

0;

Mismatches

1;

Indels Length

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76

97.4%; 99.4%;

Score 1640.4; Pred. No. 0;

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1677; 9. Gaps

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CC CDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA csequences, which can be used in vaccines. The polynucleotide sequences cc can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and cd diagnosis of diseases associated with inappropriate secretory cc protein, membrane protein expression. The nucleic acids and complementary cc sequences may also be used as DNA probes in diagnostic assays complementary cc e.g. polymerase chain reactions (PCR) to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane cc polypeptides and their role in metabolism. The polypeptides may be used ca antigens in the production of antibodies against them and in assays to identify modulators (agonists and antagonists) of expression and cativity. The antibodies and antagonists of expression and continued in a samples (e.g. by enzyme linked immunosorbant assay continued as diagnostic agents for detecting the presence of the cathories and dishates of as therapeutic apents for detecting the presence of the cathories and dishates of antibodies and include rheumatoid arthuries and dishates of antibodies may be treated include rheumatoid arthuries and dishates of antibodies which may be treated include rheumatoid arthuries and dishates of antibodies and include rheumatoid arthuries and dishates and antibodies and antibodies and antibodies and antib
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11-JAN-2000;
02-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID 323; 609pp + CD ROM; English.
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          Novel neokine
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Disclosure; Fig 1; 123pp; English.

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cysteines. The neckines have a variety of functions, including modulation of signal transduction, inhibition of angiogenesis, regulation of inflammation and inhibition of chemoattraction. Neckines may therefore be useful in the treatment of various proliferative disorders or diseases: several types of cancer, inflammation, psoriasis, and immune rejection following skin graft and kidney transplantation. Neckines, their associated polynucleotides, homologues and antibodies can be used in screening for drugs or compounds which modulate their activity and therefore may be important for the development of new therapeutics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primers were designed to amplify the region between these contigs. These primers were designed to amplify the region between these contigs. These primers were hil2/227f (AAX08966) and hil2/526r (AAX08967).

To obtain a full open reading frame (ORF), the 5' end of the single contig was generated via the RACE (rapid amplification of cDNA ends) procedure, using primer til2racel (AAX08968). The neokines are a novel family of chemoattractant cytokines (chemokines) and are ligands for previously identified putative G protein-coupled receptor termed RDC1 (also called the neokine receptor). The neokines are members of the non-ELR-CXC subfamily of chemokines characterised by the absence of an ELR motif and the presence of a CXC signature motif. The CXC motif encompasses 4 highly conserved cysteine residues, with the first two conserved that the conserved amino acid residue, heaver.
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Sequence 1564 BP; 462 A; 384 C; 334 G; 384 T; 0 other;

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cal Similarity 98.3%; Pred. No. 1e-289; 1482; Conservative 0; Mismatches 0; Indels ccacgccgcgcccctccggtcagcatgaggctcctggcggcgcgcgc	Qy	B 8		Дb	Qy	ф	Qy	Db	Qy	B	Qy	Db	Qy	Db	Qy	Db	Qy	Best Matc	Tand
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767		708	Db
838	ggggaaaactcatgcctttcctttttaaaaaatgcttttttgtatttgtccatacgtcac	779	Qy
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subfamily of chemokines characterised by the absence of an ELR motif and the presence of a CXC signature motif. The CXC motif encompasses 4 highly conserved cysteine residues, with the first two cysteines separated by one non-conserved amino acid residue; however, the neckines show some atypical features which distinguish them from previously characterised CXC chemokines e.g., the presence of approximately 5 residues between the third and fourth conserved cysteines. The neckines have a variety of functions, including modulation of signal transduction, inhibition of angiogenesis, regulation of inflammation and inhibition of chemoattraction. Neckines may therefore be useful in the treatment of various proliferative discrete a discrete control that the discrete control of the co
                                       inflammation, psoriasis, and immune rejection following skin graft and kidney transplantation. Neokines, their associated polynucleotides, homologues and antibodies can be used in screening for drugs or compounds which modulate their activity and therefore may be important for the development of new therapeutics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence represents macaque neokine-1 cDNA. The neokines are a novel family of chemoattractant cytokines (chemokines) and are ligands for a previously identified putative G protein-coupled receptor termed RDC1 (also called the neokine receptor). The neokines are members of the non-ELR-CXC
Sequence
                                                                                                                                                                         proliferative disorders or diseases: several types of cancer,
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P-PSDB; AAY31615.
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                   This invention describes novel human nucleic acid (cDNA) sequences (A), that are highly expressed in uterine tumour tissue and which have anticancer and cytostatic activity. (A) are used (i) for recombinant expression of polypeptides (B) and (ii) to isolate complete genes. (B) are used (i) to identify agents suitable for treatment of uterine or endometrial cancer; (ii) directly for treating these forms of cancer
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generation of specific antibodies. (A) are identified by assembling ESTS (expressed sequence tags) from a particular tissue type before comparison of expression patterns. This allows a significantly longer fragment of the gene to be revealed, so should reduce the number of failures associated with the fact that ESTs from different libraries may represent different parts of the same unknown gene, distorting the estimated frequency of occurrence in a particular tissue. AAZ41981-Z42121 represent EST fragments derived from a human endometrium tumour cDNA library which
                                                encode the protein sequences represented in AAY59941-Y60328
Sequence 1630 BP; 477 A; 392 C; 327 G; 434 T; 0 other;
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                                                                                                 This invention describes novel polypeptides and their encoding nucleic acids derived from human pancreatic tumor tissue which have cytostatic activity. The sequences are also useful in producing pharmaceutical compositions for treatment of pancreatic tumors. AAZ52858-Z53014 represent expressed sequence tag (EST) fragments derived from a human pancreatic tumor cDNA library and which encode the proteins represented in AAY73814-Y74252.
                                                                                 Sequence 1962 BP; 547
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes novel polypeptides and their encoding nucleic acids derived from human pancreatic tumor tissue which have cytostatic activity. The sequences are also useful in producing pharmaceutical compositions for treatment of pancreatic tumors. AAZ52858-Z53014 represent expressed sequence tag (EST) fragments derived from a human pancreatic tumor CDNA library and which encode the proteins represented
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                       tttggttgccagaaaaatgtgcttcattcccccctggttaatttttacacacccctaggaa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1442
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                                                                                                                                                                                               Mus sp.
                                                                                                                                                                                                                   anti-inflammatory;
                                                                                                                                                                                                                                                                                         Full-length cDNA encoding muKS1,
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 Novel polynucleotides including wounds and
                              WPI; 2000-072177/06
P-PSDB; AAY76115.
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09-NOV-1998;
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            treatment
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Claim 1; Page 210; 235pp; English.

cancer and neurological diseases. The proteins may be used to stimulate the growth and motility of keratinocytes, to inhibit the growth of cancer cells, to modulate angiogenesis and tumour vascularisation, to modulate skin inflammation, to modulate epithelial cell growth and to inhibit binding of HIV-1 to leukocytes. The invention may also be used to treat growth and developmental defects, skin wounds and hair follicle disorders. Sequences AAZ61606-Z61832 represent cDNA sequences derived from several mouse, rat or human skin cell types. Sequences AAZ61649, AAZ61725-Z61765, AAZ61802-Z61811 and AAZ61826 encode proteins with an v-terminal signal sequence, indicating that the proteins are secreted. Sequences AAZ61650-Z61668, AAZ61766-Z61780, AAZ61812-Z61817 and AAZ61827-Z61829 encode proteins with one or more putative and mouse embryonic skin, keratinocyte stem cells and transit amplifying cells. Polypeptides of the invention may be used to treat inflammation, The invention relates to novel nucleic acid sequences derived from rat dermal papilla, human keratinocytes and neonatal foreskin fibroblasts, transmembrane domains.

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Sequence 1663 BP; 404 A; 476 C; 428 ç; 355 T; 0 other;

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QΥ B Вþ 밁 Qy Вþ γ δã DЬ B Qy 밁 Ωy B δÃ B Qγ Ъ Qy 망 Qy В Qγ В Q Db Qγ QΥ DЬ Qy Db Db Ş 1535 1391 1138 1078 1018 994 754 635 tatg tatg cca¢cccatgccacacactgcccctcattagaggccgcactgtatggctgtg-tatctgc gctaagatgaccatgcgccctt----tcctctgtacatatacccttaagaacgcccccatafatttgtgattccccatgtaattcttcaatgttaaacagtgcagtcctctttcgaaa 1480 ata gtt#ggacgatcctgcgccctgccctctcctgtgtacatattgccttcagtacccctccc catgtgaggttcccaacactgtcagcaaa------aaccttaggagaaaacttaaaa | 99aBacatttccaagatcctgtgatggcgagacaaatgatccttaaagaaggtgtgggggt 1197 caagtttggttgccagaaaaatgtgcttcattcccccctggttaatttttacacaccccta 1137 cccbcccccaaagctttgtttctttccctcaccccaacagaaagtgcact----atghaggattagatgcaggacacacagccagagctacacatcctcttggcaatgggagct catpcaaagtccccagcagtgtgagaaaatgattctcgatgttcgggaggacaagggaag לננ ctt-tcccaacctgaggatttctgaaaggttcacaggttcaatatttaatgcttcagaag gcapcccagctcattcccctgtcacgaccaggcaagtgatccttaaaggagctgggtctt c-agtotacttaccagaaggagottcatccccacccc-----acccccaccc 1099 gtggcccctgttgcagggcaattgggagcaggtccttctacatctgtgcctagaggaact 1053 9t9gcccctgtagccgggcaagcaggagcaggtctctctctgcatctgttctctgaggaact 1077 tgdagctgccacaggctctcctgggcttatgactggtcagagtttcagtgtgactccact cacagctgccacgggctctcctgggcttatggccggtcacagcctcagtgtgactccaca cagcactactgctccgctccaggcttacaaagcttccgctcagagagcctggcggctctg ttgcagtgttgctccattcctagcttgggaagcttccgcttagaggtcctggcctcgg ccacacatctgagctttataagcgcctgggaggaacagtgagcatggttgagaccgttca ctatacatctgagctttataagcgcccgggaggaacaatgagcttggtggacacatttca atdtetettgtgeteeagaetteateaeaggetgettttt----cggggggacctcaagcctttcctttttaaaataaggggttttgtatttgtccatatgtca agggaaaactcatgcctttcctttttaaaaaatgctttttgtatttgtccatacgtca agqtcgcctgcgcaccagacttcattacaagtggctttttgctggggggttggcggttggcggggg acq Chitgcaaactgagggtttctgaaaggtcggctgctttggtagaagatgcttctgagg -aaccttcaaagcatgtttctttccctcaccacaacagaacatgcagtactaaagca -agggtcagtttttacattttatagctgtgtgtgaaaggcttccagatgtgagatcc ccccctcagtgaatacgcaacagcactgttctctgagttaggat 1435 1390 1420 . 1159 1364 1219 1017 993 933 693

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC99756 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tumours, modulating skin inflammation, stimulating the growth of epithelial cells, inhibiting the binding of human immunodeficiency virus (HIV)-1 to leukocytes, and treating inflammatory disease, cancer and neurological diseases. The polynucleotide can be used as a marker, in the identification of genetic disorders, and for the design of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mammalian skin cells. The polypeptide is useful for stimulating keratinocyte growth and motility, inhibiting the growth of cancer cells modulating angiogenesis, inhibiting angiogenesis and vascularisation of the control of the cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated polynucleotide used in the identification of genetic disorders and encoding polypeptides used for treating inflammatory disease, cancer and neurological diseases -
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                                                           cctccggccagcatgaggctcctggcggccgcgctgctcctgctgctcctggcgctgtgc
                                                                               cctccggtcagcatgaggctcctggcggccgcgctgctcctgctgctgctgctggcgctgtac
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                          keratinocyte growth and motility, inhibiting the growth of cancer cells, modulating angiogenesis, inhibiting angiogenesis and vascularisation of tumours, modulating skin inflammation, stimulating the growth of epithelial cells, inhibiting the binding of human immunodeficiency virus (HIV)-1 to leukocytes, and treating inflammmatory disease, cancer and
(HIV)-1 to leukocytes, and treating inflammmatory disneurological diseases. The polynucleotide can be used the identification of genetic disorders, and for the
                                                                             The present polynucleotide encodes a polypeptide which is expressed mammalian skin cells. The polypeptide is useful for stimulating keratinocyte growth and motility, inhibiting the growth of cancer or
                                                                                                                                                            New isolated polynucleotide used in the identification of disorders and encoding polypeptides used for treating infidisease, cancer and neurological diseases -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptor). The neckines are members of the non-ELR-CXC subfamily of chemokines characterised by the absence of an ELR motif and the presence of a CXC signature motif. The CXC motif encompasses 4 highly conserved cysteine residues, with the first two cysteines separated by one non-conserved amino acid residue; however, the neckines show some atypical features which distinguish them from previously characterised CXC chemokines e.g., the presence of approximately 5 residues between the third and fourth conserved cysteines. The neckines have a variety of functions, including modulation of signal transduction, inhibition of anglogenesis, regulation of inflammation and inhibition of chemoattraction. Neckines may therefore be useful in the treatment of various
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 are a novel family of chemoattractant cytokines (chemokines) and are ligands for a previously identified putative G protein-coupled receptor termed RDC1 (also called the neokine receptor). The neokines are members of the non-ELR-CXC
                                                                                                                                                                                                                                                                                                                                                                        proliferative disorders or diseases: several types of cancer, inflammation, psoriasis, and immune rejection following skin graft and kidney transplantation. Neokines, their associated polynucleotides, homologues and antibodies can be used in screening for drugs or compounds which modulate their activity and therefore may be important for the development of new therapeutics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents murine neokine-1 cDNA. The neokines
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        Claim 1;
                        Novel neokine polypeptides and polynucleotides used for regulating and treating proliferative disorders and diseases
                                                       P-PSDB;
                                                                                    Barnes TM
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Best Local :
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Pred. No. 1.2e-115;
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RESULT 15
AAF94062
ID AAF940
XX
AC AAF940
XT
DT 23-MAY
XX
Primer
XX
KW Human;
KW Theuman
XX
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OS Synthe

(first

entry)

AAF94062; 23-MAY-2001

AAF94062 standard;

DNA;

Synthetic

Human; secretory protein; membrane protein; vaccine; gene therapy; rheumatoid arthritis; diabetes; PCR primer; ss. Primer specific for DNA encoding secretory/membrane protein SEQ ID Вþ Qy 망 õ Вþ QΥ В Q

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CR AAF93917 - AAF94295 and AAF62235 - AAF62235 which are used to isolate the

CR AAF93917 - AAF94295 and AAF62235 - AAF62235 which are used to isolate the

CR CAAF93917 - AAF94295 and AAF62235 - AAF62235 which are used to isolate the

CR CAAF93917 - AAF94295 and AAF62235 - AAF62235 which are used to isolate the

CR CAAF93917 - AAF94295 and AAF62235 - AAF62235 which are used to isolate the

CR CAAF93917 - AAF94295 and AAF62235 which are used to isolate the

CR CAAF93917 - AAF94295 and AAF62235 which are primers

CR CAAF93917 - AAF94295 and AAF62235 which are proteins, and cDNA

CR CAAF93917 - AAF94295 and AAF62235 which are proteins, and cDNA

CR CAAF93917 - AAF94295 and the reaction of interprepariate sequences and the

CR CAAF93917 - AAF94295 and AAF62235 which may polymucleotide sequences

CR CAAF93917 - AAF94295 and AAF62235 which may be used as complementary

CR CAAF93917 - AAF94295 and AAF62235 which may be used as therapeutic

CR CAAF93917 - AAF94295 and AAF62235 which may be used as therapeutic

CR CAAF93917 - AAF94295 and AAF62235 which may be treated include rheumatoid

CR CAAF93917 - AAF94295 and AAF62235 which may be treated include rheumatoid

CR CAAF93917 - AAF94295 and AAF62235 which may be treated include rheumatoid

CR CAAF93917 - AAF94295 and AAF62235 which may be treated include rheumatoid

CR CAAF93917 - AAF94295 and AAF62235 which may be treated include rheumatoid
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11-JAN-2000; 2000JP-0118775.
02-MAY-2000; 2000JP-0183766.
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Pred. No. 8.6e-111;
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Result
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                    Score
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	3, Application US/09188930A 6150502 Watson, James D. Strachan, Lorna Sleeman, Matthew Onrust, Rene Murison, James Greg Murison, James Greg Murison, James Greg ENCE: 11000.1011c1 PLICATION UNMEER: US/09/188,930A LING DATE: 1998-11-09 EGO ID NOS: 348 BastSEQ for Windows Version 3.0 63 Mouse -263	3.1 1364 2 US-08-965-688-3 Sequence 3, Appli 3.1 1364 4 US-09-260-173-3 Sequence 3, Appli 3.1 1364 4 US-09-260-173-3 Sequence 3, Appli 3.0 1525 4 US-09-157-603-4 Sequence 4, Appli 3.0 1525 4 US-09-914-969-14 Sequence 4, Appli 3.0 3871 2 US-08-599-4558-3 Sequence 14, Appli 3.0 3871 2 US-08-798-181-3 Sequence 3, Appli 3.0 3871 4 US-08-728-259A-10 Sequence 3, Appli 3.0 1001 1 US-08-73-486-10 Sequence 10, Appli 3.0 1001 2 US-08-433-944-1 Sequence 11, Appli 3.0 1242 4 US-08-431-974-1 Sequence 1, Appli 3.0 1242 4 US-08-437-388-1 Sequence 1, Appli 3.0 1242 4 US-08-174-739A-1 Sequence 1, Appli 3.0 1242 4 US-08-174-739A-1 Sequence 1, Appli 3.0 1242 4 US-08-174-739A-1 Sequence 1, Appli 3.0 1254 3 US-08-51-136C-7 Sequence 1, Appli 3.0 1254 3 US-08-296-624-1 Sequence 2, Appli 3.0 134 3 US-09-248-335-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Ni, Jian
                                                           TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                     APPLICATION NUMBER: US 60/013,653
FILING DATE: 19-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Chemokine Alpha 2 NUMBER OF SEQUENCES: 10
                                                                                           TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
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              SEQUENCE CHARACTERISTICS:
LENGTH: 461 base pairs
TYPE: nucleic acid
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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TYPE: nucleic
STRANDEDNESS:
                                                                                                                         REGISTRATION NUMBER: 36, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/0 FILING DATE: 19-MAR-1997
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Li, Haodong
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Suite 600
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GENERAL INFORMATION:
APPLICANT: Watson, James D.
APPLICANT: Strachan, Lorna
APPLICANT: Strachan, Lorna
APPLICANT: Onrust, Rene
APPLICANT: Murison, James Greg
TITLE OF INVENTION: Compositions Isolated From Sk:
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1011c1
CURRENT APPLICATION MUMBER: US/09/188,930A
CURRENT FILING DATE: 1998-11-09
NUMBER OF SEQ ID NOS: 348
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Best Local :
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Patent No. 6150502
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127..375
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Pred. No. 1.8e-89;
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RESULT 4
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                               GENERAL INFORMATION:
APPLICANT: Watson, James D.
                                                                Sequence 270, Application US/09188930A Patent No. 6150502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Fas
SEQ ID NO 38
LENGTH: 766
 APPLICANT:
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Best Local Similarity 79.5%;
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ORGANISM: mouse
                                                                                                                                                            694 ccagctcgcctgcgcaccagacttcattacaagtggcttttt 735
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Strachan, Lorna
Sleeman, Matthew
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NUMBER OF SEQ ID NOS: 348
SOFTWARE: FBSTSEQ for Windows Version 3.0
SEQ ID NO 272
LENGTH: 234
TYPE: DNA
GRGANISM: Mouse
US-09-188-930-272
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 270
LENGTH: 288
TYPE: DNA
ORGANISM: Mouse
US-09-188-930-270
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APPLICANT: Watson, James D.
APPLICANT: Strachan, Lorna
APPLICANT: Sleeman, Matthew
APPLICANT: Sleeman, Matthew
APPLICANT: Murison, James Greg
TITLE OF INVENTION: Compositions Isolated From Skin Cells
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1011c1
CURRENT APPLICATION NUMBER: US/09/188,930A
CURRENT FILLING DATE: 1998-11-09
                                                                         Best Local Similarity 100.0%;
Matches 234; Conservative
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APPLICANT: Murison, James Greg
TITLE OF INVENTION: Compositions Isolated From
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1011c1
CURRENT APPLICATION NUMBER: US/09/188,930A
CCURRENT FILING DATE: 1998-11-09
241 aagtggtacaacgcctggaacgagaagcgcagggtctacgaagaatag
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                                                                                               Score 234; DB 3;
Pred. No. 3.8e-44;
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                                                                               Indels
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; ORGANISM: Mouse
US-09-188-930-271
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APPLICANT: Murison, James Greg
APPLICANT: Murison, James Greg
TITLE OF INVENTION: Compositions Isolated From Skin Cells
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1011c1
CURRENT APPLICATION NUMBER: US/09/188,930A
CURRENT FILING DATE: 1998-11-09
NUMBER OF SEQ ID NOS: 348
SOFTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 271
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                                                                                                                                                                                 Sequence 14, Application US/08232463 Patent No. 5670367
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                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH:
                                                                                APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKNER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
                                                  CORRESPONDENCE ADDRESS:
                                                                        NUMBER OF SEQUENCES:
                                                                                                                                              APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                         389 gtgtccaggtaccgaggtcaggagcactgcctgcaccccaagctgcagagcaccaagcgc 448
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                61 gaaatgaagccaaagtacccacactgcgaggagaagatggttatcgtcaccaccaagagc 120
             STREET:
               ADDRESSEE: Foley & Lardner STREET: 1800 Diagonal Road, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gaaatgaagccaaagtacccgcactgcgaggagaagatggttatcatcaccaccaagagc 388
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Alexandria
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Sleeman, Matthew
Onrust, Rene
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                                                                                                                                                DORNER, F.
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94.9%;
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Pred. No. 8.1e-40;
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US-08-232-463-14 PTZgpt-F1s
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NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
TELEFAX: (703)683-4109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: line
IMMEDIATE SOURCE:
CLONE: pTZgpt-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                             1000 AAAAATAGTTATAGTAGCCGCACTCGATGGG 970
                                                                                                                                                                                                                                                                      442 caagcgcttcatcaagtggtacaacgcctggaacgagaagcgcagggtctacgaagaata 501
                                                                                                                                                                                                                                                                                                                                                                                             322 gaagctggaaatgaagccaaagtacccgcactgcgaggagaagatggttatcatcaccac 381
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 22313-0299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: BENT, Stephen A. REGISTRATION NUMBER: 29,768 REFERENCE/DOCKET NUMBER: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                            gggtgaaaaacctcagaagggaaaactccaaaccagttgggagacttgtgcaaaggactt 561
                                                                                                                                                                                                                                                                                                         AGCTCCCTCGACCTGCAGCCAAGCTCGGAATTAATTCTGTGAGCGTATGGCAAACGAAGG 1001
                                                                                     ttctcacaggcataagacacaaattatattgttatgaagcactttttaccaacggtca 681
                         gtttttacattttatagctgcgtgcgaaagg 712
                                                                                                                       40;
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GENERAL INFORMATION:
APPLICANT: SZOSTAK, JACK W.
APPLICANT: SZOSTAK, JACK W.
APPLICANT: COBERTS, RICHARD W.
APPLICANT: LIU, RINE
TITLE OF INVENTION: SELECTION OF PROTEINS USING RNA-PROTEIN
TITLE OF INVENTION: FUSIONS
FILE REFERENCE: 00786/350007
CURRENT APPLICATION NUMBER: US/09/244,796
CURRENT APPLICATION NUMBER: 60/035,963
EARLIER APPLICATION NUMBER: 60/035,963
EARLIER APPLICATION NUMBER: 60/064,491
EARLIER APPLICATION NUMBER: 60/064,491
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; LOCATION: (1)...(289)
; OTHER INFORMATION: n = A,T,C or G
US-09-007-005-17
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APPLICANT: Roberts, Richard W.
APPLICANT: Liu, Rihe
TITLE OF INVENTION: SELECTION OF PROTEINS USING RNA-PROTEIN
TITLE OF INVENTION: FUSIONS
FILE REFERENCE: 00786/350003
CURRENT APPLICATION NUMBER: US/09/007,005B
CURRENT APPLICATION NUMBER: US/09/007,005B
CURRENT FILING DATE: 1998-01-14
CURRENT FILING DATE: 1998-01-14
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EARLIER FILING DATE: 1997-11-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EARLIER APPLICATION NUMBER: 60/035,963
EARLIER FILING DATE: 1997-01-27
EARLIER APPLICATION NUMBER: 60/064,491
EARLIER FILING DATE: 1997-11-06
NUMBER OF SEO ID NOS: 33
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RESULT 10
US-08-378-313-24
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; NAME/KEY: misc_feature
; LOCATION: (1)...(289)
; OTHER INFORMATION: n = A,T,C or G
US-09-244-796-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No. 6207881
GENERAL INFORMATION:
APPLICANT: THEOLO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24 Application US/08378313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EARLIER APPLICATION NUMBER: 09/007,005
EARLIER FILING DATE: 1998-01-14
NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17
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Best Local
                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                  ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/862,493
                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: CONTROL OF FRUIT RIPENING THROUGH
TITLE OF INVENTION: GENETIC CONTROL OF ACC SYNTHASE SYNTHESIS
RUMBER OF SEQUENCES: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
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                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                         STREET: 755 Page
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        266 aaa¢aaaaaaaaaaaaaaaaa
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                                                                                                                    FILING DATE:
                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                 ZIP:
                                                                                                                                                                                                                                                                                                  COUNTRY
                                                                                                                                                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
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Similarity 13.7%;
MURASHIGE, KATE H.
                                                                                                                                                                                                                                                                                                                  California
                                                                                                                                                                                                                                                                                                                                                      755 Page Mill Road
                                                                                                                                                                                                                                                                                                  USA
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                                                                                                                                    US/08/378,313
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Pred. No. 0.00081;
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US-08-378-313-24
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US-08-520-678A-29
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Best Local S
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                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Rice, Charles M.
APPLICANT: ROlythalov, Alexander A.
APPLICANT: KOlythalov, Alexander A.
TITLE OF INVENTION: NOVEL 3' TERMINAL SEQUENCE OF HEPATITIS
TITLE OF INVENTION: C VIRUS GENOME AND DIAGNOSTIC AND THERAPEUTIC USES THEREOF
NUMBER OF SEQUENCES: 39
CORRESPONDENCE ADDRESS:
ADDRESSEE: Howell & Haferkamp, L.C.
STREET: 7733 Forsyth Blvd., Suite 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29,757
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 856-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 260 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
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                MOLECULE TYPE:
                                                                                                                                                                                                                                                          CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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                                                                                                                                                                                                        NAME: Henderson, Melodie W
REGISTRATION NUMBER: 37,848
REFERENCE/DOCKET NUMBER: 60
                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: St. Louis
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                                 TOPOLOGY:
                                                   STRANDEDNESS:
                                                                                                                                                        TELEFAX:
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                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                                 single
                DNA (genomic)
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72.0%;
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ER: 29190-20002.20
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                                                                                                                  ; MOLECULE TYPE: US-08-897-126-29
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Best Local Similarity
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                                                                    Query Match
                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 260 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Rice, Charles M.
APPLICANT: Kolykhalov, Alexander A.
APPLICANT: Kolykhalov, Alexander A.
TITLE OF INVENTION: NOVEL 3' TERMINAL SEQUENCE OF HEPATITIS
TITLE OF INVENTION: C VIRUS GENOME AND DIAGNOSTIC AND THERAPEUTIC USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 314-727-5188
                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       492 acgaagaatagggtgaaaaacctcagaagggaaaactccaaaccagttgggagacttgtg 551
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492 acgaagaatagggtgaaaaacctcagaagggaaaactccaaaaccagttgggaagacttgtg 551
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Howell & Haferkamp, L.C. STREET: 7733 Forsyth Blvd., Suite 1400
                                                                                                                                                                                                                                                TELEPHONE: 314 - 727 - 6092
                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 37,848
REFERENCE/DOCKET NUMBER: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE:
                                                                                                                                                                                    TYPE:
                                                                                                                                                                                                                                                                                                                          NAME: Henderson, Melodie W. REGISTRATION NUMBER: 37,848
                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/897,126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE:
                                                         Local Similarity
                                                                                                                                                     TOPOLOGY: linear
                                                                                                                                                                   STRANDEDNESS:
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                                                                                                                                                                                nucleic acid
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                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Louis
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                                                                                                                                    DNA (genomic)
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                                                          Score 54.8; DB 4;
Pred. No. 0.00096;
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                                             Mismatches
                                                                            DB 4;
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US-08-504-459-5
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 Sequence 25,
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                               1774 AAAAAAAAAAAAAAAAAAAAAA 1798
                                                                                                            589 aaaaaaaaaaaaaaaaaa 613
                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 512/418-3000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Adhesin Genes and Proteins involved TITLE OF INVENTION: Trichomonas Vaginalis Cytoadherence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                  Local Similarity
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Relace-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               132 AA 131
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Wilson, Mark B.
REGISTRATION NUMBER: 37,259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/504,459
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Application US/09248335
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                                                                                                                                                                                                                                                                                                                                                    1804 base pairs
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WENTION: Adhesin Genes and Proteins Involved
                                                                                                                                                                      Conservative
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/desc = "DNA"
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                                                                                                                                                                             Score 54.6; DB 2; Length 1804; Pred. No. 0.0023;
                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17,
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EARLIER FILING DATE: 1997-September-05
NUMBER OF SEQ ID NOS: 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/248,335
CURRENT FILING DATE: 1999-02-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: O'KEEF, DANIEL
TITLE OF INVENTION: PLANT GLUTATHIONE-S-TRANSFERASE ENZYMES
FILE REFERENCE: CL-1128-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: MCGONIGLE, APPLICANT: O'KEEF, DA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Osman Ph.D., Richard Aron
REGISTRATION NUMBER: 36,627
                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                                                                                                                                                   MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: ORIGIN OF REPLICATION COMPLEX GENES NUMBER OF SEQUENCES: 24
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   REFERENCE/DOCKET NUMBER:
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4 Embarcadero Center, Suite 3400
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GAVIN, Kimberly
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KOBAYASHI, Ryuji
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Pred. No. 0.0038;
0; Mismatches 46; Indels 0
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TELECOMMUNICATION INFORMATION:

**TELEPHONE: (415) 494-8700

**TELEFAX: (415) 494-8771

**TELEFAX: (415) 494-8771

**TELEFAX: (415) 494-8771

**INFORMATION FOR SEQ ID NO: 17:

**SEQUENCE CHARACTERISTICS:

**LENGTH: 3214 base pairs

**TRANDEDNESS: double

**TOPOLOGY: Linear

**MOLECULE TYPE: CDNA

**FEATURE:

**POPOLOGY: Linear

**MOLECULE TYPE: CDNA

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FEATURES SOURCE ORGANISM COMMENT REFERENCE VERSION KEYWORDS RESULT BG678414 ACCESSION DEFINITION AUTHORS TITLE JOURNAL source Hcmo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 976) NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC) Plate: LLAM10604 row: k co High quality sequence stop: Tissue Procurement: James Cleaver, M.D. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov 642625062F1 NCI_CGAP_Skn4 Homo sapiens cDNA clone IMAGE:4750214 5', BG678414.1 GI:13909811 mRNA sequence. BG678414 DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can LLAM10604 row: k column: 15 /Organism="Homo sapiens" /db_xref="taxon:9606" /clone="IMAGE:4750214" /clone_lib="NCI_CGAP_Skn4" /tissue_type="squamous cell carcinoma" /lab_host="DH10B (T1 phage-resistant)" 1. .976 Location/Qualifiers 976 bp TIRNA 01-MAY-2001 be

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97.7%;
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Pred. No. 1.3e-110;
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BG696197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contract: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: James Cleaver, M.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA
CDNA Library Arrayed Genomics, Inc.
Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
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plate: LLAM10696 row: j column:
High quality sequence stop: 852.
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National Institutes of Health, Mammalian Gene Collection (MGC)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/Clone_lib="NCI_CGAP_Skn3"
/lab_host="DH10B (T1 phage-resistant)"
/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: skin; vector: pCMV-SPORT6; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally. Primer: Oligo dT
Average insert size 1.5kb. Library constructed by Life
Technologies. Note: this is a NCI_CGAP Library."

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                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
Email: cgapbs r@mail.nih.gov
Tissue Procurement: James Cleaver, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM10604 row: i column: 16
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National Institutes of Health, Mammalian Gene Collection (MGC)
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Location/Qualifiers
/tissue_type="squamous cell carcinoma"
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/clone_lib="NCI_CGAP_Skn4"
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Site_2: Sall; Cloned unidirectionally. Primer: Oligo dr.
Average insert size 1.5kb. Library constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
a 198 c 173 g 237 t
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99.6%;
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caccctaggaaacatttccaagatcctgtgatggcgagacaaatgatccttaaagaaggt
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Tissue Procurement: James Cleaver, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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Plate: LLAM10609 row: j column: 09
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National Institutes of Health, Mammalian Gene Collection (MGC)
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/lab_host="DH10B (T1 phage=resistant);
/note="organ: skin; Vector: pcW-SPORT6; Site_1: Not1;
/note="organization: skin; Vector: skin; Vec
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/clone_lib="NCI_CGAP_Skn4"
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/db_xref="taxon:9606"
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Pred. No. 4.9e-104;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Li,W.B., Gruber,C., Jessee,J. and Polayes,D. Full-length cDNA libraries and normalization Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 845)
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                                                                          231
                                                                                                                                                                                                                                                                                                                                                                                                                                   91006 EVRY cedex - France segref@genoscope.cns.fr.
                                                                                                           cloned into the Not I and Eco RV sites of the pcWSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies. Contact: Feng Liang Life Technologies, a division of Invitrogen 9800 Medical Center Drive Rockville, Maryland 20850, USA Fax: (1) 301 610 8371 Email: fliang@lifetech.com URL:
                                                                                                                                                                                                                            /clone_lib="LTI_NFL006_PL2"
/tissue_type="placenta"
/note="Vector: pCMVSPORT 6; Site_1: NotI; lst strand cDNA
was primed with a NotI-oligo(dT) primer. Five prime end
enriched, double-stranded cDNA was digested with Not I and
                                                                                            http://fulllength.invitrogen.com"
                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
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Score 797.4;
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Query Match

47.3%;

Gene Collection (MGC)

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                    EST.
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              1134 cctaggaaacatttccaagatcctgtgatggcgagacaaatgatccttaaagaaggtgtg 1193
                                                                                    1074 aact¢aagtttggttgccagaaaaatgtgcttcattcccccctggttaatttttacacac 1133
                                                                                                                                                       481 CCTA¢GAAACATTTCCAAGATCCTGTGATGGCGAGACAAATGATCCTTAAAGAAGGTGTG
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Plate: LLAM10600 row: h column: 12
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Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: James Cleaver, M.D.
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National Institutes of Health, Mämmalian
Unpublished (1999)
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 794)
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/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: skin; Vector: pCMV-SPORT6; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.5kb. Library constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
a 191 c 167 g 226 t
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/clone="IMAGE:4748603"
/clone_lib="NCI_CGAP_Skn4"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TCCC
                                                         831;
                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: James Cleaver, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
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BG679890
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National Institutes of Health, Mammalian Gene Collection
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1 (bases 1 to 860)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                              found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                   Plate: LLAM10607 row: i column: 24
                                                                     Similarity
                                                                                                                                                                                                                                                                                                                   quality sequence stop: 858.
                                                         Conservative
                                                                                                                                    /tissue_type="squamous cell carcinoma"
/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: skin; Vector: pCMV-SPORT6; Site_1: Not1;
Site_2: SalI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.5kb. Library constructed by Life Technologies. Note: this is a NCI_CGAP Library."
1 205 c 180 g 244 t
                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                     /clone="IMAGE:4751327"
/clone_lib="NCI_CGAP_Skn4"
                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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i1384 tttccctcaccacaacagaacatgcagtactaaagcaatatatt-tgtgattccccatgt 1442
                                                                              1324 gcaatacacagctacagacacacattctgttgacaagggaaaaccttcaaagcatgtttc 1383
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                                                                                                                                                                                                                                                                                    1204 caacctgaggatttctgaaaggttcacaggtttcaatatttaatgcttcagaagcatgtga 1263
                                                   481 GCAATACACAGCTACAGACACACTTCTGTTGACAAGGGAAAACCTTCAAAGCATGTTTC 540
                                                                                                                                                    421 GGTTCCCAACACTGTCAGCAAAAACCTTAGGAGAAAACTTAAAAATATATGAATACATGC 480
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/db_xref="taxon:9606"
/clone="IMAGE:4592520"
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/clone="IMAGE:4592520"
/clone="IMAGE:4592520"
/clone="IMAGE:4592520"
/clone="IMAGE:4592520"
/lab_bost="DHIUB (TI phage-resistant)"
/note="Organ: kidney; Vector: pDNR-LIB (Clontech); Site_1:
Sfil (ggccgctcggc); Site_2: Sfil (ggccattatggcc); 5' and
3' adaptors were used in cloning as follows: 5' adaptor
sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence:
5'-ATTCTAGAGGCCGACGACGACTG-dT(30)BM-3' (where B = A,
C, or G and N = A, C, G, or T). Average insert size 1.65
kb (range 0.5-4.0 kb). 15/15 colonies contained inserts
by PCR. This library was enriched for full-length clones
and "see Conservations of the colonies contained inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and was constructed by Clontech Laboratories (Palo Alto, CA). Note: this is a NIH_MGC Library."

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685 TTTATAAGCGCCCGGGAGGAACAATGAGCTTGGTGAACACATTTCATWGCAGTGTTGCTC
                                                                                                     792 gcctttcctttttaaaaaatgcttttttgtatttgtccatacgtcactatacatctgagc 851
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: segref@genoscope.cns.fr,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genoscope - Centre National de Sequencage
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                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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/clone_ib="LTI_NFL006_PL2"
/tissue_type="placenta"
/tissue_type="placenta"
/note="Vector: pCMVSPORT 6; Site_1: NotI; 1st strand cDNA
/note="Vector: pCMVSPORT 6; Site_1: NotI; 1st strand cDNA
was primed with a NotI-oligo(dT) primer. Five prime end
enriched, double-stranded cDNA was digested with Not I and
cloned into the NotI and Eco RV sites of the pCMVSPORT 6
vector. Library was normalized. Library was constructed by
Life Technologies. Contact: Feng Liang Life Technologies,
a division of Invitrogen 9800 Medical Center Drive
Rockville, Maryland 20850, USA Fax: (1) 301 610 8371
Email: fliang@llfetech.com URL:
http://ciii/nach.instructed.com/URL;
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97.4%;
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                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: James Cleaver, M.D.
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                     BG681049 756 bp mRNA
602628871F1 NCI_CGAP_Skn4 Homo sapiens
                                                                                                                                                                                                 NIH-MGC http://mgc.nci.nih.gov/.
national Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                    http://image.llnl.gov
Plate: LLAM10613 row: n column: 18
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quality sequence stop: 752.
Location/Qualifiers
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              agaacatgcagtactaaagcaatatatttgtgattccc 1437
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/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: skin; Vector: pCMV-SPORT6; Site_1: Not1;
/note="Organ: skin; Vector: pCMV-SPORT6; Site_2: Sal1; Cloned unidirectionally. Primer: Oligo dT.
Site_2: Sal1; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.5kb. Library constructed by Life Technologies. Note: this is a NCI_CGAP Library."

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/clone_lib="NCI_CGAP_Skn4"
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Adachi, J., Aizawa, K., Akahira, S., Akimura, T., Aono, H., Arai, A., Arakawa, T., Carninci, P., Fukuda, S., Fukunishi, Y., Furuno, M., Haragaki, T., Hara, A., Hayatsu, N., Hiramoto, K., Hiraoka, T., Hori, F., Imotani, K., Ishi, Y., Itoh, M., Izawa, M., Kato, H., Kawai, J., Kojima, Y., Konno, H., Kouda, M., Koya, S., Kurihara, C., Matsuyama, T., Miyazaki, A., Nishi, K., Nomura, K., Numazaki, R., Ohno, M., Okazaki, Y., Okido, T., Owa, C., Saito, H., Saito, R., Sakai, C., Sakai, K., Sano, H., Sasaki, D., Shibata, K., Shibata, Y., Shinagawa, A., Shiraki, T., Sasaki, D., Shibata, K., Shibata, Y., Tagawa, A., Shiraki, F., Tanamaura, T., Yasunishi, A., Yasunishi, 
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                                                                                                                                                                                                                                                                                                                                                                     Submitted (10-JUL-2000) Yoshihide Hayashizaki, The Institute of Physical and Chemical Research (RIKEN), Laboratory for Genome Exploration Research Group, RIKEN Gemomic Sciences Center (GSC), RIKEN Yokohama Institute; 1-7-22 Suchiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan (E-mail:genome-res@gsc.riken.go.jp, URL.http://genome.gsc.riken.go.jp/, Tel:81-45-503-9222, Fax:81-45-503-9216)
                                                                                                                                                                                                                                                                                                             Please visit our web site (http://genome.gsc.riken.go.jp/) for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yoshida, K., Yoshino, M., Muramatsu, M. and Hayashizaki, Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nature 409, 685-690 (2001) 5 (bases 1 to 1668)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itch,M., Konno,H., Okazaki,Y., Muramatsu,M. and Hayyashizaki,Y. Normalization and subtraction of cap-trapper-selected cDNAs to prepare full-length cDNA libraries for rapid discovery of new genes Genome research. 10 (10), 1617-1630 (2000)
                                                                                                                                                                                                                                                                                           further details.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Functional annotation of a full-length mouse cDNA collection
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High-efficiency full-length cDNA cloning
Methods in enzymology. 303, 19-44 (1999)
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/db_xref="GI:12835906"
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/clone_lib="RIKEN formula"/
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/db_xref="MGD:MGI:1897974"
/db_xref="MGD:MGI:1915056"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits,
cDNA Library Preparation: Michael J.
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Plate: LLAM10709 row:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gctgtacaccgcgcgtgtggacgggtccaaatgcaagtgctcccgggaaggggacccaagat 303
                                                                                                                                                                                               CDNA Library Preparation: CLONTECH Laboratories, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLCM1573 row: f column: 11
                                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONTECH Laboratories, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BG57207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              602592543F1 NIH_MGC_79 Homo sapiens cDNA clone IMAGE:4719754 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BG572071.1 GI:13579724
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (bases 1 to 858)
                                                                                                                                                                  quality sequence stop: 635
Location/Qualifiers
/clone="IMAGE:4719754"
/clone_lib="NIH_MGC_79"
/clone_lib="NIH_MGC_79"
/lab_host="DH10B (TI phage-resistant)"
/note="Organ: placenta; Vector: pDNR-LIB (Clontech);
Site_1: SfiI (99ccgcctcggcc); Site_2: SfiI (9gccattatggcc
                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                      . 858
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RESULT 1
BG619277
                                              SOURCE
                                                               KEYWORDS
                                                                                  VERSION
                                                                                                         ACCESSION
                                                                                                                                           DEFINITION
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                       ORGANISM
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                                                                                                                                                                                                                                                    1516 tac¢ettaagaacgeeeeeteeaeaeaetgeeeeeeagtatatgeegeattgtaetgetg 1575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1456 taa#cagtgcagtcctctttcgaaagctaagatgaccatgcgccctttcctctgtacata 1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1396 caa¢agaacatgcagtactaaagcaatatatttgtgattccccatgtaattcttcaatgt 1455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1276 tgtþagcaaaaaccttaggagaaaacttaaaaatatatgaatacatgcgcaatacacagc 1335
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1156 ctgkgatggcgagacaaatgatccttaaagaaggtgtgggggtctttcccaacctgaggat 1215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1096 aaaltgtgcttcattcccccctggttaatttttacacaccctaggaaacatttccaagatc 1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 481 TACCCTTAAGAACGCCCCCCCCACACACTGCCCCCCCAGTATATGCCGCATTGTACTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             421 TAAACAGTGCAGTCCTCTTTCGAAAGCTAAGATGACCATGCGCCCTTTCCTCTGTACATA 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       361 CAACAGAACATGCAGTACTAAAGCAATATATTTGTGATTCCCCCATGTAATTCTTCAATGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121 CTGFGATGGCGAGACAAATGATCCTTAAAGAAGGTGTGGGGTCTTTCCCAACCTGAGGAT 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       181 TTCFGAAAGGTTCACAGGTTCAATATTTAATGCTTCAGAAGCATGTGAGGTTCCCCAACAC
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Homo sapiens
Eukaryota; M
                                                                                                                                BG619277 707 bp mRNA
602619402F1 NIH_MGC_79 Homo
                                                               BG619277.1 GI:13670648
EST.
                                                                                                   BG61927
                                                                                                                  mRNA sequence.
                                          numan.
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|Similarity 98.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Library."
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Pred. No. 2.8e-79;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                  sapiens cDNA clone IMAGE:4733112 5',
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Вb QY 밁 Qy DЬ

Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

QΥ

γQ

뫄 QΥ Дb Qy 밁 Ş g Qy DЪ Qγ 밁 Qy Вb

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FEATURES
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 643; Conservative
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                                                                                                                                                                                                                                                                                                                                                                         1036 caagcaggagcaggtctctctgcatctgttctctgaggaactcaagtttggttgccagaa 1095
                                                                                                                                        1276 tgtcagcaaaaaccttaggagaaaacttaaaaatattgaatacatgcgcaatacacagc 1335
                                                                                                                                                                                                                                               1156 ctgtgatggcgagacaaatgatccttaaagaaggtgtgggggtctttcccaacctgaggat 1215
             1456 taaacagtgcagtcctctttcgaaagctaagatgaccatgcgccctttcctctgtacata 1515
                                                                                                                                                                                                                                                                                                               121 CTGTGATGGCGAGACAAATGATCCTTAAAGAAGGTGTGGGGTCTTTCCCCAACCTGAGGAT 180
                                                            361 CAACAGAACATGCAGTACTAAAGCAATATATTTGTGATTCCCCCATGTAATTCTTCAATGT 420
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TAAACAGTGCAGTCCTCTTTCGAAAGCTAAGATGACCATGCGCCCTTTCCTCTGTACATA 480
                                                                           caacagaacatgcagtactaaagcaatatatttgtgattccccatgtaattcttcaatgt 1455
                                                                                                                           Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONTECH Laboratories, Inc.
cDNA Library Preparation: CLONTECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop: 623.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         http://image.llnl.gov
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   36.7%;
99.1%;
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                                                                                1156 ctgtgatggcgagacaaatgatccttaaagaaggtgtgggggtctttcccaacctgaggat 1215
181 TTCTGAAAGGTTCACAGGTTCAATATTTAATGCTTCAGAAGCATGTGAGGTTCCCCAACAC
                                                                 121 CTGTGATGGCGAGACAAATGATCCTTAAAGAAGGTGTGGGGGTCTTTCCCCAACCTGAGGAT 180
                                                                                                                                61 AAATGTGCTTCATTCCCCCCTGGTTAATTTTTACACACCCTAGGAAACATTTCCAAGATC 120
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                                                                                                                                                                                                                                                                   618;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens cDNA cdA clones Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yang,Y., Song,H., Peng,Y., Gu,Y., Gao,G., Xiao,H., Xu,X., Li,N., Qian,B., Liu,F., Qu,J., Gao,X., Cheng,Z., Xu,Z., Zeng,L., Xu,S., Gu,W., Tu,Y., Jia,J., Fu,G., Ren,S., Zhong,M., Lu,G., Hu,R., Chen,J., Chen,Z. and Han,Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AV733062 cdA Homo sapiens cDNA clone cdAAJE12 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Zeguang Han Chinese National Human Genome Center at Shanghai Chinese National Human Genome Center at Shanghai 351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai 201203, P. R. China Tel: 86-21-50801919(ex.45) Fax: 86-21-50801922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AV733062.1 GI:10850607
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: hanzg@chgc.sh.cn
This clone is available at CHGC in Shanghai.
                                                                                                                                                                                                                                                                                                                                                                 187
                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                 /lab_host="BM25.8"
/note="Vector: pTriplEx2; Site_1: sfiIA; Site_2: sfiIB"
/note="Vector: pTriplEx2; Site_1: sfiIA; Site_2: sfiIB"
                                                                                                                                                                                                                                                                                                                                                                                                            /tissue_type="pheochromocytoma"
/dev_stage="Adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="cdAAJE12"
/clone_lib="cdA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                36.4%; Score 612.6;
98.6%; Pred. No. 1.5
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